

CONVOLVED SUBSAMPLING ESTIMATION WITH APPLICATIONS TO BLOCK BOOTSTRAP

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The block bootstrap approximates sampling distributions from dependent data by resampling data blocks. A fundamental problem is establishing its consistency for the distribution of a sample mean, as a prototypical statistic. We use a structural relationship with subsampling to characterize the bootstrap in a new and general manner. While subsampling and block bootstrap differ, the block bootstrap distribution of a sample mean equals that of a k -fold self-convolution of a subsampling distribution. Motivated by this, we provide simple necessary and sufficient conditions for a convolved subsampling estimator to produce a normal limit that matches the target of bootstrap estimation. These conditions may be linked to consistency properties of an original subsampling distribution, which are often obtainable under minimal assumptions. Through several examples, the results are shown to validate the block bootstrap for means under significantly weakened assumptions in many existing (and some new) dependence settings, which also addresses a standing conjecture of Politis, Romano and Wolf [*Subsampling* (1999) Springer]. Beyond sample means, convolved subsampling may not match the block bootstrap, but instead provides an alternative resampling estimator that may be of interest. Under minimal dependence conditions, results also broadly establish convolved subsampling for general statistics having normal limits.

1. Introduction. Subsampling and block bootstrap are two common nonparametric tools for statistical inference under dependence; see Politis, Romano and Wolf [29] and Lahiri [19], respectively, for monographs on these. Both aim to approximate distributions of statistics with correlated data, and both are data resampling methods that use blocks of neighboring observations to capture dependence. The subsampling approach of Politis and Romano [28] treats data blocks as small scale renditions of the original data, which provides replication of a statistic for estimating a sampling distribution. The block bootstrap differs philosophically by using data blocks as building material to recreate the original data. Essentially, data blocks are randomly selected and pasted together to reproduce a full-scale set of bootstrap data, as proposed by Künsch [17] and Liu and Singh [25] for extending

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Efron's [11] bootstrap to time series. As noted in Politis, Romano and Wolf [29] (cf. Section 3.9), subsampling is often valid under weak assumptions about the dependent process, basically requiring that a nondegenerate (possibly nonnormal) limit exist for the sampling distribution being approximated. In contrast, the block bootstrap applies to mean-like statistics with normal limits and typically requires comparatively much stronger assumptions for its validity. Case-by-case treatments are commonly needed to validate the bootstrap across differing dependence conditions. However, while perhaps not widely recognized, subsampling can in fact be used to verify the block bootstrap in some cases, which is a theme of this work.

We investigate estimators defined by the k -fold self-convolution of a subsampling distribution, and establish a new and general theory for their consistency to normal limits. There are two basic motivations for considering such convolved subsampling. The first is that, in the fundamental case of sample means, the block bootstrap estimator is a k -fold self-convolution of a subsampling distribution (centered and normalized), where the level k of convolution corresponds to the number of resampled blocks. This observation was originally noted by Politis, Romano and Wolf [29], who suggested this aspect as a potential technique for showing the validity of the bootstrap. Specifically, they conjectured that convolved subsampling might provide a route for establishing the block bootstrap under minimal conditions for nonstationary, strongly mixing processes, in analogy to bootstrap results existing for stationary, mixing series due to Radulović [30, 31]. For the bootstrap under dependence, the findings for the sample mean in [30, 31] have stood out as an exception, verifying the method under the same weak assumptions as subsampling (i.e., conditions essentially needed for a normal limit to exist). By investigating the convolved subsampling approach here, we can answer the above conjecture affirmatively. Moreover, we show convolved subsampling leads to a simple and unified procedure for establishing the block bootstrap for sample means under further types of processes and much weaker conditions than previously considered, such as linear time processes, long-memory sequences, (nonstationary) almost periodic time series and spatial fields. Hence, convolved subsampling estimation allows for bootstrap consistency under dependence to be generally extended under the same weak assumptions used by subsampling, containing the conclusions of Radulović [30, 31] for stationary time series as a special case.

While connections to the bootstrap are useful, our study of convolved subsampling estimation is intended to be broad, applying also to general statistics with normal limits and with arbitrary levels of convolution. Consistency results often do not require particular assumptions about the underlying dependent process, but are rather formulated in terms of mild convergence properties of the original subsampling distribution and its variance. Furthermore, we show that a consistent subsampling variance is not only sufficient, but essentially necessary, for the consistency of convolved subsampling (and the block bootstrap in some cases). Due to its importance, we also provide tools for verifying the consistency of subsampling variance estimators.

For general statistics beyond the sample mean, the convolved subsampling distribution may differ from the block bootstrap, which relates to a second motivation for our development. That is, a general theory for convolved subsampling may be of interest in its own right, as the approach can be computationally less demanding than the block bootstrap while also potentially enhancing ordinary subsampling for approximating sampling distributions with normal limits. In fact, there has been recent interest in establishing generalized types of subsampling estimation for complicated statistics under various dependence structures, where numerical studies suggest such methods can exhibit better finite sample performance than standard subsampling when the target distribution is normal; for example, see Lenart [22] and Sharipov, Tewes and Wendler [32] for spectral estimates and U-statistics, respectively, with time series. While not formally recognized as such, however, these proposed methods are exactly convolved subsampling estimators. By exploiting this realization, our results can facilitate future work and allow such previous findings with generalized subsampling to be demonstrated in an alternative, simpler manner with weaker assumptions; see Section 5 for illustrations of the examples mentioned above.

Section 2 describes convolved subsampling estimation and its connection to block bootstrap. General distributional results for convolved subsampling are given in Section 3, while Section 4 presents some applications with differing dependence structures. Section 4.1 provides a broad result for convolved subsampling estimation with statistics from mixing time series. Under weak conditions, Sections 4.2–4.5 apply convolved subsampling for demonstrating the block bootstrap for sample means with nonstationary time series (Section 4.2 and the conjecture of Politis, Romano and Wolf [29]), linear time processes (Section 4.3), long-range dependence (Section 4.4) and spatial data (Section 4.5). Section 5 describes relationships to other recent work with generalized subsampling, and Section 6 provides a short treatment of independent data. A numerical study of subsampling, block bootstrap and convolved subsampling appears in Section 7, while Section 8 contains concluding remarks. The proofs of main results are given in the Supplementary Material [15].

Finally, to be clear, we stress that a central advantage of classical subsampling is its validity for nonnormal limits (cf. Section 4.4), which convolved subsampling does not share. The convolution of a subsampling distribution essentially induces a sum of independently resampled terms so that, like the block bootstrap, reproducing a nonnormal limit is impossible. However, for approximating normal targets, convolved subsampling does inherit the applicability of subsampling under weak conditions with general statistics.

2. Description of convolved subsampling estimators.

2.1. *Problem background and original subsampling estimation.* Consider data X_1, \dots, X_n from a real-valued process equipped with a probability structure P .

For concreteness, we may view such observations as arising from a time series process $\{X_t\}$, though spatial and other data schemes may be treated as well. Based on X_1, \dots, X_n , consider the problem of approximating the distribution of

$$T_n \equiv \tau_n(t_n(X_1, \dots, X_n) - t(P)),$$

involving an estimator $t_n \equiv t_n(X_1, \dots, X_n)$ of a parameter $t(P)$ and a sequence of positive scaling factors τ_n yielding a distributional limit for T_n . For example, if $t_n(X_1, \dots, X_n) \equiv \bar{X}_n = \sum_{i=1}^n X_i/n$ is the sample mean, then $t(P)$ may correspond to a common process mean μ and T_n may be defined with usual scaling $\tau_n = \sqrt{n}$ under weak time dependence. Denote the sampling distribution function of T_n as $F_n(x) = P(T_n \leq x)$, $x \in \mathbb{R}$.

We next define the subsampling estimator of F_n ; see [28]. For a positive integer $b \equiv b_n < n$, let $\{(X_i, \dots, X_{i+b-1}) : i = 1, \dots, N_n\}$ denote the set of $N_n \equiv n - b + 1$ overlapping data blocks, or subsamples, of length b . To keep blocks relatively small, the block size is often assumed to satisfy $b^{-1} + b/n + \tau_b/\tau_n \rightarrow 0$ as $n \rightarrow \infty$. For each subsample, we compute the statistic as $t_{n,b,i} = t_b(X_i, \dots, X_{i+b-1})$ and define a “scale b ” version of $T_n \equiv \tau_n(t_n(X_1, \dots, X_n) - t(P))$ as $\tau_b[t_{n,b,i} - t_n]$ for $i = 1, \dots, N_n$. Letting $I(\cdot)$ denote the indicator function, the subsampling estimator of F_n is given by

$$(2.1) \quad S_{n,\text{SUB}}(x) = \frac{1}{N_n} \sum_{i=1}^{N_n} I(\tau_b[t_{n,b,i} - t_n] \leq x), \quad x \in \mathbb{R},$$

or the empirical distribution of subsample analogs $\{\tau_b[t_{n,b,i} - t_n]\}_{i=1}^{N_n}$ (cf. [29]).

Suppose that $S_{n,\text{SUB}}$ is consistent for the distribution of T_n , which has an asymptotically normal $N(0, \sigma^2)$ limit for some $\sigma > 0$, that is, as $n \rightarrow \infty$,

$$(2.2) \quad T_n \xrightarrow{d} N(0, \sigma^2),$$

$$(2.3) \quad \sup_{x \in \mathbb{R}} |S_{n,\text{SUB}}(x) - \Phi(x/\sigma)| \xrightarrow{P} 0,$$

where $\Phi(\cdot)$ is the standard normal distribution function. We wish to consider estimators of the distribution F_n of T_n formed by self-convolutions of the subsampling estimator $S_{n,\text{SUB}}$. This provides a general class of block resampling estimators in its own right, but also has explicit connections to block bootstrap estimators in the important case that the statistic of interest $t_n(X_1, \dots, X_n) = \bar{X}_n$ is a sample mean, as described next.

2.2. *Convolved subsampling and connections to block bootstrap.* Let k_n be a sequence of positive integers and define a triangular array $\{Y_{n,1}^*, \dots, Y_{n,k_n}^*\}_{n \geq 1}$, where, for each n , $\{Y_{n,j}^*\}_{j=1}^{k_n}$ are i.i.d. variables following the subsampling distribution $S_{n,\text{SUB}}$, as determined by (2.1) from data X_1, \dots, X_n . For $n \geq 1$, define a

centered and scaled sum

$$(2.4) \quad Z_n^* \equiv \frac{1}{\sqrt{k_n}} \sum_{j=1}^{k_n} (Y_{n,j}^* - m_{n,\text{SUB}}),$$

where $m_{n,\text{SUB}} \equiv \int x dS_{n,\text{SUB}}(x) = N_n^{-1} \sum_{i=1}^{N_n} \tau_b[t_{n,b,i} - t_n]$ is the mean of the subsampling distribution $S_{n,\text{SUB}}$, and let

$$C_{n,k_n}(x) \equiv P_*(Z_n^* \leq x), \quad x \in \mathbb{R},$$

denote the induced resampling distribution P_* of Z_n^* . Then C_{n,k_n} represents the k_n -fold self-convolution of the subsampling distribution $S_{n,\text{SUB}}$, with appropriate centering/scaling adjustments. That is,

$$C_{n,k_n}(x) = \underbrace{S_{n,\text{SUB}} * S_{n,\text{SUB}} * \cdots * S_{n,\text{SUB}}}_{k_n \text{ times}}(x\sqrt{k_n} + k_n m_{n,\text{SUB}}), \quad x \in \mathbb{R}.$$

We consider C_{n,k_n} as an estimator of the distribution F_n of T_n and formulate general conditions under which such convolved subsampling is consistent.

As suggested earlier, such results have direct implications for block bootstrap estimation as well, because the convolved subsampling estimator C_{n,k_n} exactly matches a block bootstrap estimator in the basic sample mean case $t_n(X_1, \dots, X_n) = \bar{X}_n$. To illustrate, consider approximating the distribution of $T_n = \sqrt{n}(\bar{X}_n - \mu)$ where $t(P) \equiv \mu = E\bar{X}_n$ and $\tau_n = \sqrt{n}$. In this setting, the block bootstrap uses an analog

$$(2.5) \quad T_n^* = \sqrt{n_1}(\bar{X}_{n_1}^* - E_*\bar{X}_{n_1}^*)$$

based on the average $\bar{X}_{n_1}^* \equiv n_1^{-1} \sum_{i=1}^{n_1} X_i^*$ from a block bootstrap sample $X_1^*, \dots, X_{n_1}^*$ of size $n_1 \equiv k_n b$, which is defined by drawing k_n blocks of length b , independently and with replacement, from the subsample collection $\{(X_i, \dots, X_{i+b-1}) : i = 1, \dots, N_n\}$ and pasting these together (where above $E_*\bar{X}_{n_1}^* = N_n^{-1} \sum_{i=1}^{N_n} b^{-1} \times \sum_{j=i}^{i+b-1} X_j$ denotes the bootstrap expectation of $\bar{X}_{n_1}^*$); see Chapter 2, Lahiri [19]. Most typically, the number of resampled blocks is taken as $k_n = \lfloor n/b \rfloor \rightarrow \infty$ so that the bootstrap sample recreates the approximate length $\lfloor n/b \rfloor b \approx n$ of the original sample. The bootstrap distribution of T_n^* here is then equivalent to the convolved subsampling distribution C_{n,k_n} . This is because T_n^* has the same resampling distribution as Z_n^* in (2.4) as a sum of k_n i.i.d. block averages $(Y_{n,i}^* - m_{n,\text{SUB}})/\sqrt{k_n}$, with each $Y_{n,i}^*$ drawn from $S_{n,\text{SUB}}$ in (2.1) where $t_n = \bar{X}_n$ and $\tau_b[t_{n,b,i} - t_n] = \sqrt{b}[b^{-1} \sum_{j=i}^{i+b-1} X_j - \bar{X}_n]$, $1 \leq i \leq N_n$, for the sample mean case. Consequently, if convolved subsampling estimators C_{n,k_n} are shown to be valid under weak conditions, such results entail that block bootstrap estimation is as well. In the following, we make comprehensive use of the fact that C_{n,k_n} is always and exactly a block bootstrap estimator whenever the underlying statistic $t_n(X_1, \dots, X_n) = \bar{X}_n$ is a sample mean; this holds true across all the various dependent data structures considered here, including cases where the usual block bootstrap from (2.5) requires modification for sample means (cf. long-range dependence in Section 4.3).

3. Fundamental results for convolved subsampling. From (2.1) and the subsampling mean $m_{n,\text{SUB}} \equiv \int x dS_{n,\text{SUB}}(x) = N_n^{-1} \sum_{j=1}^{N_n} \tau_b[t_{n,b,i} - t_n]$, we have the variance of the original subsampling distribution $S_{n,\text{SUB}}$ as

$$\hat{\sigma}_{n,\text{SUB}}^2 \equiv \int (x - m_{n,\text{SUB}})^2 dS_{n,\text{SUB}}(x) = \frac{1}{N_n} \sum_{j=1}^{N_n} (\tau_b[t_{n,b,i} - t_n] - m_{n,\text{SUB}})^2,$$

which estimates the asymptotic variance σ^2 of T_n as in (2.2) (cf. [29]). Note that $\hat{\sigma}_{n,\text{SUB}}^2$ is also the variance of the convolved subsampling distribution C_{n,k_n} [i.e., the variance of the i.i.d. sum from (2.4)]. Correspondingly, $\hat{\sigma}_{n,\text{SUB}}^2$ is then a block bootstrap variance estimator when applied to sample means.

Sections 3.1–3.3 provide basic distributional results for convolved subsampling estimators, describing when and how these have normal limits. These findings do not involve particular assumptions about the process $\{X_t\}$, but are instead expressed through properties of the original subsampling distribution $S_{n,\text{SUB}}$ and, specifically, convergence of the subsampling variance $\hat{\sigma}_{n,\text{SUB}}^2$. Such subsampling properties can often be verified under weak assumptions about a process, allowing the limit behavior of convolved estimators C_{n,k_n} , and the block bootstrap, to be established under minimal conditions. Results in Section 3.1 address the important case where the original subsampling distribution $S_{n,\text{SUB}}$ has a normal limit (2.3), as is often natural when the statistic $T_n \xrightarrow{d} N(0, \sigma^2)$ is asymptotically normal. These findings are expected to be the most practical for establishing convolved subsampling C_{n,k_n} estimation with normal targets (2.2). Dropping the condition that $S_{n,\text{SUB}}$ converges to a normal law but assuming convolved estimators C_{n,k_n} are based on increasing convolution $k_n \rightarrow \infty$ of $S_{n,\text{SUB}}$, Section 3.2 characterizes the convergence of C_{n,k_n} to normal limits through the subsampling variance $\hat{\sigma}_{n,\text{SUB}}^2$. In many problems involving the block bootstrap for sample means (cf. Section 4), where T_n has a normal limit (2.2), these results provide both necessary and sufficient conditions for the validity of the block bootstrap as well as convolved subsampling generally. Finally, because convergence $\hat{\sigma}_{n,\text{SUB}}^2 \xrightarrow{p} \sigma^2$ of the subsampling variance emerges as central to the behavior of convolved estimators C_{n,k_n} , Section 3.3 develops basic results for establishing this feature.

3.1. *Convolution of subsampling distributions with normal limits.* Theorem 1 provides a sufficient condition for the general validity of the convolved estimator C_{n,k_n} via fundamental subsampling quantities, $S_{n,\text{SUB}}$ and $\hat{\sigma}_{n,\text{SUB}}^2$.

THEOREM 1. *Suppose (2.3) holds [i.e., $\sup_{x \in \mathbb{R}} |S_{n,\text{SUB}}(x) - \Phi(x/\sigma)| \xrightarrow{p} 0$] and $\hat{\sigma}_{n,\text{SUB}}^2 \xrightarrow{p} \sigma^2 > 0$ as $n \rightarrow \infty$. Then*

$$\sup_{x \in \mathbb{R}} |C_{n,k_n}(x) - \Phi(x/\sigma)| \xrightarrow{p} 0 \quad \text{as } n \rightarrow \infty$$

for any positive integer sequence k_n .

Furthermore, when (2.2) holds additionally [i.e., $T_n \xrightarrow{d} N(0, \sigma^2)$], then C_{n,k_n} is consistent for the distribution F_n of T_n ,

$$\sup_{x \in \mathbb{R}} |C_{n,k_n}(x) - F_n(x)| \xrightarrow{P} 0 \quad \text{as } n \rightarrow \infty.$$

To reiterate, the integer sequence $k_n, n \geq 1$, need not even be convergent in Theorem 1. The consistency of the subsampling variance estimator $\hat{\sigma}_{n,\text{SUB}}^2$ automatically guarantees that, for any amount k_n of convolution of $S_{n,\text{SUB}}$, the convolved subsampling estimator C_{n,k_n} will have a normal limit if the subsampling distribution $S_{n,\text{SUB}}$ does. In other words, if (2.2)–(2.3) hold so that $S_{n,\text{SUB}}$ is consistent, then C_{n,k_n} will be as well provided $\hat{\sigma}_{n,\text{SUB}}^2 \xrightarrow{P} \sigma^2$. When the statistic $t_n(X_1, \dots, X_n) = \bar{X}_n$ is a sample mean, then C_{n,k_n} again denotes a block bootstrap estimator based on k_n resampled blocks, which is thereby consistent under Theorem 1 for any sequence k_n , including the common choice $k_n = \lfloor n/b \rfloor \rightarrow \infty$.

Proposition 1 next characterizes the convolved subsampling estimator C_{n,k_n} under bounded levels k_n of convolution. In this case, a normal limit for the subsampling estimator $S_{n,\text{SUB}}$ entails the same for the convolved estimator C_{n,k_n} , provided the mean $m_{n,\text{SUB}} \equiv \int x dS_{n,\text{SUB}}(x)$ of the subsampling distribution converges to zero. But, if the subsampling mean $m_{n,\text{SUB}}$ converges in this fashion, a normal limit for C_{n,k_n} with bounded $\{k_n\}$ is equivalent to a normal limit for the original subsampling distribution $S_{n,\text{SUB}}$.

PROPOSITION 1. *Suppose $\sup_n k_n < \infty$:*

(i) *If (2.3) holds [i.e., $\sup_{x \in \mathbb{R}} |S_{n,\text{SUB}}(x) - \Phi(x/\sigma)| \xrightarrow{P} 0$], then*

$$\sup_{x \in \mathbb{R}} |C_{n,k_n}(x) - \Phi(x/\sigma)| \xrightarrow{P} 0 \quad \text{as } n \rightarrow \infty$$

if and only if $m_{n,\text{SUB}} \equiv \int x dS_{n,\text{SUB}}(x) \xrightarrow{P} 0$.

(ii) *If $m_{n,\text{SUB}} \xrightarrow{P} 0$ as $n \rightarrow \infty$, then (2.3) holds if and only if*

$$\sup_{x \in \mathbb{R}} |C_{n,k_n}(x) - \Phi(x/\sigma)| \xrightarrow{P} 0 \quad \text{as } n \rightarrow \infty.$$

When the original subsampling estimator $S_{n,\text{SUB}}$ is consistent for a distribution with a normal limit [i.e., (2.2)–(2.3)], both Theorem 1 and Proposition 1 show that the convolved subsampling estimator C_{n,k_n} is consistent under an additional subsampling moment condition. With bounded levels k_n of convolution, the additional condition under Proposition 1 is that the subsampling mean converge $m_{n,\text{SUB}} \xrightarrow{P} 0$. But, for general and potentially unbounded k_n , the additional condition from Theorem 1 for consistency of C_{n,k_n} is a convergent subsampling variance $\hat{\sigma}_{n,\text{SUB}}^2 \xrightarrow{P} \sigma^2$. With diverging amounts $k_n \rightarrow \infty$ of convolution, which is often encountered in

practice and in connection to the block bootstrap, it turns out that convergence $\hat{\sigma}_{n,\text{SUB}}^2 \xrightarrow{P} \sigma^2$ is also *necessary* for consistency of the convolved estimator C_{n,k_n} , as treated in the next section.

3.2. *Unbounded convolution of subsampling distributions.* We next consider the behavior of convolved subsampling estimators with unbounded convolution $k_n \rightarrow \infty$ as $n \rightarrow \infty$, which arises, for example, with the block bootstrap C_{n,k_n} for sample means with $k_n = \lfloor n/b \rfloor$ resampled blocks. Results here do not explicitly require convergence of the original subsampling estimator $S_{n,\text{SUB}}$ to a normal limit (2.3). While a reasonable condition in problems where the target quantity $T_n \xrightarrow{d} N(0, \sigma^2)$ is asymptotically normal, limits for $S_{n,\text{SUB}}$ are not directly necessary for convolved estimators C_{n,k_n} to yield normal limits from increasing convolution k_n of $S_{n,\text{SUB}}$. However, convergence of the subsampling variance $\hat{\sigma}_{n,\text{SUB}}^2$ is crucial, as shown next.

THEOREM 2. *Suppose $k_n \rightarrow \infty$ and $\int_{|x| \geq \sqrt{k_n}\epsilon} x^2 dS_{n,\text{SUB}}(x) \xrightarrow{P} 0$ for each $\epsilon > 0$ as $n \rightarrow \infty$:*

(i) *Then*

$$\sup_{x \in \mathbb{R}} |C_{n,k_n}(x) - \Phi(x/\sigma)| \xrightarrow{P} 0$$

if and only if $\hat{\sigma}_{n,\text{SUB}}^2 \xrightarrow{P} \sigma^2 > 0$ as $n \rightarrow \infty$.

(ii) *When $\hat{\sigma}_{n,\text{SUB}}^2 \xrightarrow{P} \sigma^2 > 0$ as $n \rightarrow \infty$, then C_{n,k_n} is a consistent estimator of the distribution F_n of T_n if and only if $T_n \xrightarrow{d} N(0, \sigma^2)$ [i.e., a normal limit (2.2) for T_n holds or $\sup_{x \in \mathbb{R}} |F_n(x) - \Phi(x/\sigma)| \rightarrow 0$].*

For an *unbounded* sequence $k_n \rightarrow \infty$ of convolution (e.g., block bootstrap with $k_n = \lfloor n/b \rfloor$ concatenated blocks), Theorem 2 imposes no direct assumption on the convergence of the original subsampling distribution, but rather that $S_{n,\text{SUB}}$ fulfills a mild truncated second moment property. From this, the convergence of the convolved subsampling estimator C_{n,k_n} to a normal limit is completely determined by the subsampling variance $\hat{\sigma}_{n,\text{SUB}}^2$ under Theorem 2. Furthermore, when $\hat{\sigma}_{n,\text{SUB}}^2$ converges, the convolved estimator C_{n,k_n} will be valid for estimating the distribution F_n of a target quantity T_n having a normal limit [Theorem 2(ii)]. In cases where T_n fails to have a normal limit, the convolved estimator C_{n,k_n} does not apply.

The following corollary of Theorem 2 shows that a convolved estimator C_{n,k_n} will quite generally have a normal limit, provided that the subsampling variance converges $\hat{\sigma}_{n,\text{SUB}}^2 \xrightarrow{P} \sigma^2 > 0$ and that some other basic feature exists for the subsampling distribution $S_{n,\text{SUB}}$ or for composite statistics $\{\tau_b[t_{n,b,i} - t_n] \equiv$

$\tau_b[t_b(X_i, \dots, X_{i-b+1}) - t_n(X_1, \dots, X_n)]\}_{i=1}^{N_n \equiv n-b+1}$ defining $S_{n,\text{SUB}}$ in (2.1). Essentially, Corollary 1 entails that the truncated second moment assumption in Theorem 2 is mild in conjunction with $\hat{\sigma}_{n,\text{SUB}}^2 \xrightarrow{P} \sigma^2$.

COROLLARY 1. *Suppose one of the following conditions (C.1)–(C.4) holds:*

(C.1) *for some distribution J_0 with variance $\sigma^2 > 0$, $S_{n,\text{SUB}}(x) \xrightarrow{P} J_0(x)$ as $n \rightarrow \infty$ for any continuity point $x \in \mathbb{R}$ of J_0 ;*

(C.2) *for some $\epsilon_0 > 0$, $N_n^{-1} \sum_{i=1}^{N_n} [\tau_b(t_{n,b,i} - t_n)]^{2+\epsilon_0} = O_p(1)$;*

(C.3) *the subsample-based sequence $\{T_{b,i}^2 \equiv \tau_b^2[t_{n,b,i} - t(P)]^2 : i = 1, \dots, N_n\}_{n \geq 1}$ is uniformly integrable and $T_n \equiv \tau_n(t_n - t(P)) = O_p(\tau_n/\tau_b)$;*

(C.4) *$\{X_i\}$ is stationary, $\{T_n^2 : n \geq 1\}$ is uniformly integrable, and $\tau_b/\tau_n = O(1)$.*

Then, as $n \rightarrow \infty$,

$$\sup_{x \in \mathbb{R}} |C_{n,k_n}(x) - \Phi(x/\sigma)| \xrightarrow{P} 0$$

for any sequence k_n with $\lim_{n \rightarrow \infty} k_n = \infty$ if and only if $\hat{\sigma}_{n,\text{SUB}}^2 \xrightarrow{P} \sigma^2 > 0$.

REMARK 1. For reference, note $\tau_b/\tau_n \rightarrow 0$ often holds with subsample scaling as $n \rightarrow \infty$ so that conditions $\tau_b/\tau_n = O(1)$ and $T_n = O_p(\tau_n/\tau_b)$ are mild.

Hence, if $k_n \rightarrow \infty$ and $\hat{\sigma}_{n,\text{SUB}}^2 \xrightarrow{P} \sigma^2$, then the convolved estimator C_{n,k_n} will converge to a normal limit if the subsampling distribution $S_{n,\text{SUB}}$ is convergent (C.1) or has an appropriate stochastically bounded moment (C.2), or if the subsampling statistics related to computing $S_{n,\text{SUB}}$ have uniformly integrable second moments (C.3)–(C.4). Condition (C.4) is a special case of (C.3) under stationarity, and corresponds to an underlying assumption of Radulović [30, 31] for examining the block bootstrap estimator C_{n,k_n} of a sample mean with stationary, mixing processes; see also Remark 2 to follow. When restricted to Condition (C.1), the “ \Leftarrow ” part of Corollary 1 corresponds to an initial convolved subsampling result due to Politis, Romano and Wolf [29] (Proposition 4.4.1) for unbounded convolution $k_n \rightarrow \infty$, which was developed for establishing the block bootstrap estimator C_{n,k_n} for the sample mean of nonstationary data, as reconsidered here in Section 4.2. Note that, for inference with T_n having a normal $N(0, \sigma^2)$ limit (2.2), Condition C.1 in Corollary 1 is perhaps most natural and approachable by verifying convergence $S_{n,\text{SUB}}$ to a normal (2.3). In which case, the implication of Corollary 1 (involving $k_n \rightarrow \infty$) for guaranteeing that convolved subsampling and block bootstrap estimators replicate normal limits when $\hat{\sigma}_{n,\text{SUB}}^2 \xrightarrow{P} \sigma^2$ also becomes a special case of Theorem 1 (involving any k_n).

REMARK 2. For block bootstrap estimation of the sample mean $T_n = \sqrt{n}(\bar{X}_n - EX_1)$ with strongly mixing, stationary processes, Radulović [30, 31] provides necessary and sufficient conditions for convergence of C_{n,k_n} (with $k_n = \lfloor n/b \rfloor \rightarrow \infty$) to a normal limit, assuming $\{T_n^2 : n \geq 1\}$ is uniformly integrable. Under such assumptions, the main result there is that normal limits for both C_{n,k_n} and T_n are equivalent. In comparison, the necessary and sufficient conditions for normality of the block bootstrap estimator C_{n,k_n} for a mean in Theorem 2 are perhaps more basic in that the conclusions of [30, 31], under the additional assumptions made there, follow from Theorem 2 (cf. Corollary 1). In this sense, Theorem 2 broadly reframes the findings in [30, 31], by not involving particular process assumptions (i.e., stationarity or mixing) and applying to convolved subsampling estimators C_{n,k_n} with general statistics and arbitrarily increasing convolution levels $k_n \rightarrow \infty$. Further connections to, and extensions of, the results of Radulović [30, 31] are made in Section 4.1 for strongly mixing processes.

3.3. *Consistency of subsampling variance estimators.* Theorems 1–2 demonstrate that the subsampling variance $\hat{\sigma}_{n,\text{SUB}}^2$ plays a key role in the convergence of the convolved subsampling estimator C_{n,k_n} generally, and of the block bootstrap for the sample mean in particular. However, convergence of the subsampling distribution $S_{n,\text{SUB}}$ itself is often much easier to directly establish under weak assumptions about the process $\{X_t\}$; see Politis, Romano and Wolf [29] and Section 4 to follow. This raises a further question considered next: if one knows that subsampling estimator $S_{n,\text{SUB}}$ is consistent (2.3) for a normal limit, then when will the subsampling variance $\hat{\sigma}_{n,\text{SUB}}^2$ be convergent as well, thereby guaranteeing (from Theorem 1) that the convolved estimator C_{n,k_n} is also consistent? As shown in Theorem 3, a general characterization is possible as well as simple sufficient conditions based on moment properties of subsample statistics (e.g., T_b^2).

For $n \geq 1$, recall $T_n \equiv \tau_n(t_n(X_1, \dots, X_n) - t(P))$ and additionally define $T_{n,i} \equiv \tau_n(t_n(X_i, \dots, X_{i+n-1}) - t(P))$ for $i \geq 1$ from the statistic applied to (X_i, \dots, X_{i+n-1}) . Based on $N_n \equiv n - b + 1$ subsample observations of length $1 \leq b \equiv b_n < n$, define a distribution function

$$(3.1) \quad D_{n,b}(x) \equiv \frac{1}{N_n} \sum_{i=1}^{N_n} P(T_{b,i} \leq x), \quad x \in \mathbb{R},$$

as an average of subsample-based probabilities.

THEOREM 3. *Suppose (2.3) and $T_n = o_p(\tau_n/\tau_b)$ as $n \rightarrow \infty$:*

(i) *Then $\hat{\sigma}_{n,\text{SUB}}^2 \xrightarrow{P} \sigma^2 > 0$ as $n \rightarrow \infty$ if and only if, for each $\epsilon > 0$,*

$$(3.2) \quad \lim_{m \rightarrow \infty} \sup_{n \geq m} P\left(\frac{1}{N_n} \sum_{i=1}^{N_n} T_{b,i}^2 I(|T_{b,i}| > m) > \epsilon\right) = 0.$$

(ii) *Additionally, (3.2) holds whenever $\{Y_b^2 : b \geq 1\}$ is uniformly integrable, where Y_b denotes a random variable with distribution $D_{n,b}$, $n \geq 1$, from (3.1) [i.e., $P(Y_b \leq x) = D_{n,b}(x)$, $x \in \mathbb{R}$]. If (2.3) and $T_n = o_p(\tau_n/\tau_b)$ hold, uniform integrability of $\{Y_b^2 : b \geq 1\}$ is equivalent to $\int x^2 dD_{n,b}(x) = N_n^{-1} \sum_{i=1}^{N_n} ET_{b,i}^2 \rightarrow \sigma^2$ as $n \rightarrow \infty$.*

(iii) *(3.2) also holds whenever $\{X_t\}$ is stationary and $\{T_b^2 : b \geq 1\}$ is uniformly integrable.*

REMARK 3. As T_n is typically tight, the assumption $T_n = o_p(\tau_n/\tau_b)$ is often satisfied by a standard condition on block length: $b \rightarrow \infty$ with $b/n + \tau_b/\tau_n \rightarrow 0$. Block conditions are not, in fact, used or required in statements of Theorems 1–3 above. However, block assumptions are usually needed to show the original subsampling estimator $S_{n,\text{SUB}}$ is convergent as in (2.3), and examples of Section 4 shall impose block length conditions for this purpose.

Theorem 3 connects convergence (2.3) of subsampling distributions $S_{n,\text{SUB}}$ to the convergence of subsampling variances $\hat{\sigma}_{n,\text{SUB}}^2$ in a way involving no further conditions on the process or statistic beyond mild types of uniform integrability. For example, with nonstationary processes $\{X_t\}$, Theorem 3(ii) converts the problem of probabilistic convergence $\hat{\sigma}_{n,\text{SUB}}^2 \xrightarrow{p} \sigma^2$ into a more approachable one of subsample-moment convergence $N_n^{-1} \sum_{i=1}^{N_n} ET_{b,i}^2 \rightarrow \sigma^2$. To frame another implication of Theorem 3, note that many inference problems with time series involve a stationary process $\{X_t\}$ and a statistic T_n with a normal limit (2.2) such that $\{T_n^2 : n \geq 1\}$, and consequently $\{T_b^2 : b \geq 1\}$, is uniformly integrable; see Remark 2. In such problems, it suffices to simply establish the consistency of the subsampling estimator $S_{n,\text{SUB}}$ (2.3) and then the consistency of subsampling variance $\hat{\sigma}_{n,\text{SUB}}^2$ follows with no further effort [by Theorem 3(iii)] along with the consistency of the convolved subsampling estimator C_{n,k_n} (by Theorem 1). Again, with sample means, C_{n,k_n} is a block bootstrap distribution and $\hat{\sigma}_{n,\text{SUB}}^2$ is a block bootstrap variance estimator, so both will be consistent in this setting by showing that $S_{n,\text{SUB}}$ is consistent. This strategy has two advantages with the block bootstrap: showing the consistency of $S_{n,\text{SUB}}$ is often an easier prospect than considering either C_{n,k_n} or $\hat{\sigma}_{n,\text{SUB}}^2$ directly, and the consistency of $S_{n,\text{SUB}}$ (and thereby the bootstrap) can typically be established under weak process assumptions.

To illustrate, Section 4 applies the basic results here for establishing the convolved subsampling estimator C_{n,k_n} , as well as the block bootstrap for sample means, under different dependence structures.

4. Applications of convolved subsampling estimation. Section 4.1 first develops consistency results for convolved subsampling estimators with strongly mixing processes and general statistics. The remaining subsections then consider convolved subsampling for the particular case of the sample mean with the goal of

generalizing and extending results for the block bootstrap across various types of dependent data, such as nonstationary mixing time processes (Section 4.2), linear time series (Section 4.3), long-range dependent processes (Section 4.4) and spatial data (Section 4.5).

Define the strong mixing coefficient of $\{X_t\}$ as $\alpha(k) = \sup_{i \in \mathbb{Z}} \{|P(A \cap B) - P(A)P(B)| : A \in \mathcal{F}_{-\infty}^i, B \in \mathcal{F}_{k+i}^\infty\}$, $k \geq 1$, where $\mathcal{F}_{-\infty}^i$ and \mathcal{F}_{k+i}^∞ respectively denote σ -algebras generated by $\{X_t : t \leq i\}$ and $\{X_t : t \geq k + i\}$ (cf. [1], Chapter 16.2). Recall $\{X_t\}$ is strongly mixing or α -mixing if $\lim_{k \rightarrow \infty} \alpha(k) = 0$.

4.1. *Convolved subsampling for general statistics under mixing.* For mixing stationary time series, Radulović [30] proved consistency of block bootstrap estimation for $T_n = \tau_n(t_n(X_1, \dots, X_n) - t(P))$ based on the sample mean $t_n(X_1, \dots, X_n) = \bar{X}_n$ with $t(P) = EX_1$ and $\tau_n = \sqrt{n}$. The assumptions made were quite weak, requiring only:

- (a1) a stationary, α -mixing process fulfilling (2.2) [i.e., $T_n \xrightarrow{d} N(0, \sigma^2)$] and block lengths $b^{-1} + b/n \rightarrow 0$ as $n \rightarrow \infty$;
- (a2) uniformly integrable $\{T_n^2 : n \geq 1\}$.

From results in Section 3 and the equivalence between the block bootstrap and the convolved subsampling estimator C_{n,k_n} for the sample mean, a different perspective is possible for the bootstrap findings in Radulović [30]. Under only assumption (a1) above, the subsampling estimator $S_{n,\text{SUB}}$ is consistent [i.e., (2.3) holds] for the asymptotically normal distribution of $T_n = \sqrt{n}(\bar{X}_n - EX_1)$ (cf. Theorem 3.2.1, [29]), implying, by Theorem 1 here, that the block bootstrap estimator C_{n,k_n} would be consistent if the subsampling variance converges $\hat{\sigma}_{n,\text{SUB}}^2 \xrightarrow{P} \sigma^2$. But, if $S_{n,\text{SUB}}$ is consistent for a normal limit by (a1), assumption (a2) then guarantees that $\hat{\sigma}_{n,\text{SUB}}^2 \xrightarrow{P} \sigma^2$ holds by Theorem 3. Furthermore, under (a2) and with $k_n = \lfloor n/b \rfloor \rightarrow \infty$ resampled blocks as in Radulović [30, 31], convergence $\hat{\sigma}_{n,\text{SUB}}^2 \xrightarrow{P} \sigma^2$ becomes even necessary here by Theorem 2. Hence, α -mixing serves to show that the original subsampling estimator $S_{n,\text{SUB}}$ is consistent; after which, uniform integrability and stationary assure both $\hat{\sigma}_{n,\text{SUB}}^2 \xrightarrow{P} \sigma^2$ and consistency of the block bootstrap estimator C_{n,k_n} by Theorems 2–3.

Under analogously weak assumptions as those of Radulović [30], Theorem 4 next provides the general consistency of convolved subsampling estimation for general statistics arising from mixing, and possibly nonstationary, time processes. When applied to a sample mean $t_n(X_1, \dots, X_n) = \bar{X}_n$, so that C_{n,k_n} is a block bootstrap estimator, this result extends those of Radulović [30] in two ways: by allowing potential nonstationarity series and by permitting arbitrary levels k_n of convolution/block resampling (rather than the single choice $k_n = \lfloor n/b \rfloor$). When the statistic $t_n(X_1, \dots, X_n)$ is not a sample mean, C_{n,k_n} may not again match the block bootstrap but can have interest as an alternative block resampling estimator (cf. Section 5).

THEOREM 4. *Let $\{X_t\}$ be a (possibly nonstationary) strongly mixing sequence. Suppose $b^{-1} + b/n + \tau_b/\tau_n \rightarrow 0$ as $n \rightarrow \infty$; $T_n = o_p(\tau_n/\tau_b)$; (3.2) holds; and that $Y_b \xrightarrow{d} N(0, \sigma^2)$ as $n \rightarrow \infty$, for some $\sigma^2 > 0$, where each random variable Y_b , $b \equiv b_n \geq 1$, has distribution function $D_{n,b}$ from (3.1). Then, as $n \rightarrow \infty$,*

$$\sup_{x \in \mathbb{R}} |S_{n,\text{SUB}}(x) - \Phi(x/\sigma)| \xrightarrow{p} 0 \quad \text{and} \quad \hat{\sigma}_{n,\text{SUB}}^2 \xrightarrow{p} \sigma^2$$

and, for any positive integer sequence k_n ,

$$\sup_{x \in \mathbb{R}} |C_{n,k_n}(x) - \Phi(x/\sigma)| \xrightarrow{p} 0.$$

Furthermore, if (2.2) additionally holds [i.e., $T_n \xrightarrow{d} N(0, \sigma^2)$], then $S_{n,\text{SUB}}$ and C_{n,k_n} (with any k_n) are consistent for the distribution F_n of T_n :

$$\sup_{x \in \mathbb{R}} |S_{n,\text{SUB}}(x) - F_n(x)| \xrightarrow{p} 0 \quad \text{and} \quad \sup_{x \in \mathbb{R}} |C_{n,k_n}(x) - F_n(x)| \xrightarrow{p} 0.$$

While providing a broad result on the validity of convolved subsampling estimation for mixing processes, Theorem 4 also expands the general subsampling results of Politis, Romano and Wolf [29] (Chapter 4.2), which focused on $S_{n,\text{SUB}}$ for mixing series, to further include consistency of the subsampling variance $\hat{\sigma}_{n,\text{SUB}}^2$. That is, when dropping (3.2), the remaining Theorem 4 assumptions are minimal and match those of Theorem 3.2.1–4.2.1 of Politis, Romano and Wolf [29] for the consistency of $S_{n,\text{SUB}}$ to a normal limit; including (3.2) in Theorem 4 is then necessary for $\hat{\sigma}_{n,\text{SUB}}^2 \xrightarrow{p} \sigma^2$ by Theorem 3 and assures convergence of C_{n,k_n} by Theorem 1.

If the process $\{X_t\}$ is actually stationary, we immediately obtain the following result.

COROLLARY 2. *Let $\{X_t\}$ be a stationary, strongly mixing sequence. Suppose also $b^{-1} + b/n + \tau_b/\tau_n \rightarrow 0$ as $n \rightarrow \infty$; that (2.2) holds; and that (3.2) holds (e.g., uniform integrability of $\{T_n^2 : n \geq 1\}$ suffices). Then, as $n \rightarrow \infty$, the convergence results of Theorem 4 hold.*

Section 5 illustrates Theorem 4 for establishing convolved subsampling with mixing time series and several general classes of statistics. These represent cases where C_{n,k_n} differs from the block bootstrap estimator.

However, Section 4.2 first provides some further refinements with mixing processes in the sample mean case, where C_{n,k_n} matches the block bootstrap.

4.2. Block bootstrap for mixing nonstationary time processes. Consider a strongly mixing, potentially nonstationary sequence $\{X_t\}$ having a common mean parameter $EX_t = \mu \in \mathbb{R}$, which is estimated by the sample mean \bar{X}_n . In this setting and under conditions where $T_n \equiv \sqrt{n}(\bar{X}_n - \mu)$ has a normal limit (2.2),

Fitzenberger [12] established the consistency of the block bootstrap for estimating the distribution of T_n . The result, however, required the existence of a $(4 + \delta)$ -moment (i.e., $\sup_t E|X_t|^{4+\delta} < \infty$ for some $\delta > 0$) along with stringent mixing conditions and restrictions on the block length $b = o(n^{1/2})$. Politis, Romano and Wolf [29] (Example 4.4.1) showed that the subsampling estimator $S_{n,\text{SUB}}$ is consistent under weaker conditions, including only a $(2 + \delta)$ -moment. For the block bootstrap with $k_n = \lfloor n/b \rfloor$ resampled blocks, Politis, Romano and Wolf [29] also proved bootstrap consistency by applying convolved subsampling in this problem, using a weaker block assumption $b = o(n)$ than Fitzenberger [12] but otherwise with same remaining strong assumptions about the process. However, [29] (Remark 4.4.4) conjectured that the block bootstrap might be established under nonstationarity using the same weak moment/mixing conditions as the subsampling estimator $S_{n,\text{SUB}}$, just as in the case of stationary mixing processes (cf. [30]). We confirm this by the following Theorem 5.

THEOREM 5. *Let $\{X_t\}$ be a sequence of (not necessarily stationary) strongly mixing random variables with common mean μ . For some $\delta > 0$, suppose that $\sup_t E|X_t|^{2+\delta} < \infty$ and $\sum_{k=1}^\infty \alpha(k)^{\delta/(2+\delta)} < \infty$. Assume also that, for some $\sigma^2 > 0$,*

$$\lim_{n \rightarrow \infty} \sup_{i \geq 1} \left| \text{Var} \left(n^{-1/2} \sum_{t=i}^{i+n-1} X_t \right) - \sigma^2 \right| = 0.$$

Then, as $n \rightarrow \infty$, $T_n = \sqrt{n}(\bar{X}_n - \mu) \xrightarrow{d} N(0, \sigma^2)$ [i.e., (2.2) holds]. Additionally, if $b^{-1} + b/n \rightarrow 0$ as $n \rightarrow \infty$, then

$$\sup_{x \in \mathbb{R}} |S_{n,\text{SUB}}(x) - \Phi(x/\sigma)| \xrightarrow{p} 0 \quad \text{and} \quad \hat{\sigma}_{n,\text{SUB}}^2 \xrightarrow{p} \sigma^2$$

and, for any positive integer sequence k_n ,

$$\sup_{x \in \mathbb{R}} |C_{n,k_n}(x) - \Phi(x/\sigma)| \xrightarrow{p} 0.$$

Hence, with any number k_n of concatenated blocks, the block bootstrap estimator C_{n,k_n} is valid for the distribution of the sample mean under mild assumptions for mixing, and possibly nonstationary processes. Note that the assumptions of Theorem 5 resemble those essentially needed for a central limit theorem (CLT) itself (cf. Theorem 16.3.5, [1]). In particular, the assumptions also match those commonly used in the stationary case for establishing the block bootstrap; see Section 3.2 of Lahiri [19]. With the same moment condition as Politis, Romano and Wolf [29] (Theorem 4.4.1), Theorem 5 additionally shows that the original subsampling estimator $S_{n,\text{SUB}}$ is consistent under nonstationarity with even weaker mixing assumptions than considered previously $\sum_{k=1}^\infty (k + 1)^2 \alpha(k)^{\delta/(8+\delta)} < \infty$. The central message of Theorem 5, however, is that the convolved subsampling

approach allows the block bootstrap estimator C_{n,k_n} for the sample mean to be established under weak conditions similarly to $S_{n,SUB}$.

Next, consider the block bootstrap in another important example of nonstationarity, involving certain periodically correlated time series. Here, the mean function $\mu(t) \equiv EX_t$ is not constant, as in Theorem 5, but rather an almost periodic function. A real-valued function f is *almost periodic* if, for every $\epsilon > 0$, there is an $n(\epsilon) \in \mathbb{N}$ such that in every interval $I_{n(\epsilon)}$ of length $n(\epsilon)$ or greater, there is an integer $p \in I_{n(\epsilon)}$ such that

$$\sup_{t \in \mathbb{Z}} |f(t + p) - f(t)| < \epsilon;$$

see [7]. For such functions, the limit $M(f) \equiv \lim_{n \rightarrow \infty} n^{-1} \sum_{i=s}^{s+n-1} f(i)$ exists and does not depend on s . Moreover, if the set $\Lambda = \{\lambda \in [0, 2\pi) : M(g_\lambda) \neq 0\}$ is finite for $g_\lambda(t) \equiv f(t)e^{-i\lambda t}$, $t \in \mathbb{Z}$ ($i = \sqrt{-1}$), then

$$(4.1) \quad \left| \frac{1}{n} \sum_{i=s}^{s+n-1} (f(i) - M(f)) \right| \leq \frac{C}{n}$$

holds for some $C > 0$ not depending on n or s by Cambanis et al. [6]. Hence, $M(f)$ represents the mean value of an almost periodic function f . A time series is called *almost periodically correlated* (APC) if its mean and autocovariance functions are almost periodic, that is, for every fixed $\tau \in \mathbb{Z}$,

$$\mu(t) = EX_t \quad \text{and} \quad \rho_\tau(t) = EX_t X_{t+\tau}$$

are almost periodic as functions of t ; see [14]. For an APC series $\{X_t\}$, a parameter of interest is then $t(P) \equiv M(\mu) = \lim_{n \rightarrow \infty} n^{-1} \sum_{i=s}^{s+n-1} \mu(i)$ as a summary of the process mean structure, which is estimated by \bar{X}_n . Synowiecki [34] showed that the block bootstrap consistently estimates the sampling distribution of $T_n = n^{1/2}(\bar{X}_n - M(\mu))$ under appropriate conditions. By applying the convolved subsampling technique, we may extend the bootstrap results of Synowiecki [34] (Corollary 3.2) by substantially weakening the assumptions made there about $(4 + \delta)$ -moments and $\sum_{k=1}^\infty k\alpha(k)^{\delta/(4+\delta)} < \infty$.

COROLLARY 3. *Let $\{X_t\}$ be an APC sequence of strongly mixing random variables such that $\sup_t E|X_t|^{2+\delta} < \infty$ and $\sum_{k=1}^\infty \alpha(k)^{\delta/(\delta+2)} < \infty$ for some $\delta > 0$, and suppose the set $\Lambda = \{\lambda \in [0, 2\pi) : M(g_\lambda) \neq 0\}$ is finite for $g_\lambda(t) \equiv \mu(t)e^{-i\lambda t}$, $t \in \mathbb{Z}$, with $\mu(t) = EX_t$. Then all conclusions of Theorem 5 hold for $T_n = n^{1/2}(\bar{X}_n - M(\mu))$ as $n \rightarrow \infty$.*

4.3. Block bootstrap for linear time processes. Based on a sample X_1, \dots, X_n , next consider inference about the mean $EX_t = \mu \in \mathbb{R}$ of a stationary time process $\{X_t\}$ prescribed as

$$(4.2) \quad X_t = \mu + \sum_{j \in \mathbb{Z}} a_j \varepsilon_{t-j}, \quad t \in \mathbb{Z},$$

in terms of i.i.d. variables $\{\varepsilon_t\}$ with mean zero and finite variance $E\varepsilon_t^2 \in (0, \infty)$ and a real-valued sequence $\{a_j\}$ of constants where $\sum_{j \in \mathbb{Z}} a_j^2 < \infty$. The linear series $\{X_t\}$ need not be mixing and, depending on constants $\{a_j\}$, can potentially exhibit either weak or strong forms of time dependence. Using the sample mean \bar{X}_n to estimate the process mean μ , suppose that

$$(4.3) \quad \lim_{n \rightarrow \infty} n^\alpha \text{Var}(\bar{X}_n) = \sigma^2$$

for some $\sigma^2 > 0$ and exponent $\alpha \in (0, 1]$ depending on the process $\{X_t\}$. When $\alpha = 1$, the sample mean’s variance decays at a rate $O(n^{-1})$ with the sample size, as typical for weakly, or short-range, dependent processes. However, when $\alpha \in (0, 1)$, the sample mean has a variance with comparatively slower decay $O(n^{-\alpha})$, which may be associated with processes exhibiting strong or long-range forms of dependence. Long-range dependent processes are commonly characterized by slowly decaying covariances involving a long-memory exponent $\alpha \in (0, 1)$, which results in less precision (4.3) for a sample mean compared to the weak dependence case [3]. Classes of strongly dependent processes that satisfy (4.2)–(4.3) include fractional Gaussian models [26] and fractional autoregressive integrated moving averages [13].

Based on (4.3), define $T_n \equiv n^{\alpha/2}(\bar{X}_n - \mu)$ in terms of scaling $\tau_n \equiv n^{\alpha/2}$. In this setting, the convolved subsampling C_{n,k_n} once again corresponds to the block bootstrap estimator based on k_n resampled blocks, but there is a wrinkle to note. Recalling from (2.5) that the bootstrap sample mean $\bar{X}_{n_1}^*$ is created from a bootstrap sample of length $n_1 = k_n b$, the bootstrap rendition of T_n here is given by

$$(4.4) \quad T_n^* \equiv b^{(1-\alpha)/2} (n_1)^{\alpha/2} (\bar{X}_{n_1}^* - E_* \bar{X}_{n_1}^*)$$

rather than the analog $T_n^* = (n_1)^{\alpha/2} (\bar{X}_{n_1}^* - E_* \bar{X}_{n_1}^*)$ of (2.5). While intuitive, the latter is incorrect under long memory and produces a degenerate bootstrap [18]. Instead, the bootstrap from (4.4) requires an adjustment $b^{(1-\alpha)/2}$, which disappears under weak dependence $\alpha = 1$ whereby bootstrap versions of T_n then match in (2.5) and (4.4). Interestingly, convolved subsampling estimator C_{n,k_n} automatically corresponds to the correct bootstrap rendition T_n^* in (4.4) under both weak $\alpha = 1$ and strong $\alpha \in (0, 1)$ dependence.

Considering the sample mean from stationary linear processes (4.2) ranging over short- or long-range dependence, Kim and Nordman [16] showed the consistency of the block bootstrap distribution C_{n,k_n} (when $k_n = \lfloor n/b \rfloor$) and bootstrap variance $\hat{\sigma}_{n,\text{SUB}}^2$. Via convolved subsampling, we may generalize their results. For linear processes $\{X_t\}$ satisfying (4.2)–(4.3), the sample mean $T_n \equiv n^{\alpha/2}(\bar{X}_n - \mu)$ has a normal limit (2.2) (cf. [8]) and the subsampling estimator $S_{n,\text{SUB}}$ is also consistent [i.e., (2.3) holds] under mild assumptions (cf. [27]). Hence, by primitively assuming (2.2)–(2.3) to hold, Corollary 4 next extends the block bootstrap to general stationary processes with sample means satisfying a variance condition (4.3), which includes results of Kim and Nordman [16] for linear processes as a special case.

COROLLARY 4. *Let $\{X_t\}$ be a stationary process with mean $\mu \in \mathbb{R}$ satisfying (4.3) for some $\alpha \in (0, 1]$, and suppose that (2.2)–(2.3) hold for $T_n \equiv n^{\alpha/2}(\bar{X}_n - \mu)$. Then, as $n \rightarrow \infty$,*

$$\hat{\sigma}_{n,\text{SUB}}^2 \xrightarrow{P} \sigma^2 \quad \text{and} \quad \sup_{x \in \mathbb{R}} |C_{n,k_n}(x) - \Phi(x/\sigma)| \xrightarrow{P} 0$$

for any positive integer sequence k_n .

Corollary 4 is an application of Theorems 1 and 3 for stationary processes which may not be strongly mixing. Our exposition has assumed the exponent $\alpha \in (0, 1]$ to be known. Upon replacing α with an estimator $\hat{\alpha} \equiv \hat{\alpha}(X_1, \dots, X_n)$ where $|\hat{\alpha} - \alpha| \log n \xrightarrow{P} 0$, the conclusions of Corollary 4 still hold; see Remark 3 of [16] for further details.

4.4. *Block bootstrap under long-range dependence.* This section briefly mentions the block bootstrap with additional types of long-memory sequences. Beyond linear processes, the sample mean of a long-range dependent sequence may converge to a nonnormal limit, such as the case for certain subordinated Gaussian processes considered by Taqqu [35] and Dobrushin and Major [10] [e.g., $X_t = G(Z_t)$ as a function G of a long-range dependent Gaussian series $\{Z_t\}$]. For such time series, Lahiri [18] proved that the block bootstrap sample mean always has a normal limit, so that the block bootstrap fails if the original sample mean is asymptotically nonnormal. This result is in concordance with our Theorem 2(ii).

Zhang et al. [36] considered subsampling for a wider class of long-memory series that includes both subordinated Gaussian processes as well types of linear processes (4.2). Namely, sequences $X_t = K(Z_t)$, $t \in \mathbb{Z}$, formed by a measurable transformation K of a long-range dependent linear process

$$Z_t = \varepsilon_t + \sum_{j=1}^{\infty} j^{-\beta} L(j) \varepsilon_{t-j}, \quad t \in \mathbb{Z},$$

defined with i.i.d. mean zero, finite variance innovations $\{\varepsilon_t\}$, an index parameter $1/2 < \beta < 1$ and slowly varying function $L(\cdot)$. They distinguish two cases, depending on β and the so-called power rank $p \geq 1$ of K . In the first case [i.e., $p(2\beta - 1) > 1$], the transformation K diminishes long-range dependence, and the sample mean converges to a normal limit. In the second case [i.e., $p(2\beta - 1) < 1$], the transformed process $X_t = K(Z_t)$ remains strongly dependent and the sample mean has a normal limit only when $p = 1$.

Assuming a constant function $L(\cdot) = C$ in the above formulation, the variance of a sample mean satisfies (4.3) [i.e., $\lim_{n \rightarrow \infty} n^\alpha \text{Var}(\bar{X}_n) = \sigma^2 > 0$] with a long-memory exponent $\alpha \equiv \min\{1, p(2\beta - 1)\} \in (0, 1]$ that changes between cases of weak $\alpha = 1$ or strong $\alpha = p(2\beta - 1) \in (0, 1)$ dependence (cf. Lemma 1, [36]). For the sample mean, Zhang et al. [36] established consistency of several subsampling

estimators as well as convergence of $\hat{\sigma}_{n,\text{SUB}}^2$. Thus, by slightly recasting results of [36] and applying our Corollary 4, we may show the validity of the block bootstrap C_{n,k_n} for estimating the distribution of $T_n \equiv n^{\alpha/2}(\bar{X}_n - \mu)$, $\mu = EX_t$, for transformed linear processes exhibiting either short- or long-range dependence. To the best of our knowledge, the bootstrap has not yet been investigated for such processes.

COROLLARY 5. *For $X_t = K(Z_t)$, $t \in \mathbb{Z}$, as above, suppose (4.3) holds for $\alpha = \min\{1, p(2\beta - 1)\} \in (0, 1]$ along with conditions of Theorem 1 in [36] [involving a block $b \propto n^a$ for some $a \in (0, 1]$] with either $p(2\beta - 1) > 1$, or $p = 1$ and $(2\beta - 1) < 1$. Then, for $T_n = n^{\alpha/2}(\bar{X}_n - \mu)$ as $n \rightarrow \infty$, both (2.2)–(2.3) hold and*

$$\hat{\sigma}_{n,\text{SUB}}^2 \xrightarrow{P} \sigma^2 \quad \text{and} \quad \sup_{x \in \mathbb{R}} |C_{n,k_n}(x) - \Phi(x/\sigma)| \xrightarrow{P} 0$$

for any positive integer sequence k_n .

As with subordinated Gaussian processes [18], consistency of the block bootstrap or convolved estimator C_{n,k_n} for the sample mean only follows in cases where a CLT holds. For subordinated Gaussian processes and statistics other than the sample mean, Betken and Wendler [4] proved the general consistency of the subsampling estimator, and Bai and Taquq [2] established weak conditions for the subsample size. When the original statistic has a normal limit, consistency of convolved subsampling will follow by our Theorem 2 by showing convergence of $\hat{\sigma}_{n,\text{SUB}}^2$ (which, as [2] and [4] consider stationary processes, can hold by Theorem 3 and uniform integrability).

4.5. Spatial data. While convolved subsampling results have been presented for processes $\{X_t\}$ indexed by time t to ease the exposition, Theorems 1–3 also apply to more general processes, including spatial random fields. In the Supplementary Material [15], we illustrate this with spatial data on a grid, for which various authors have considered block bootstrap and subsampling; see Lahiri [19] (Chapter 12) and Politis, Romano and Wolf [29] (Chapter 5) and references therein. Under appropriate assumptions for the stationary random field, the spatial sample mean has a normal limit and we establish convolved subsampling under mixing conditions from Lahiri [20] (Section 4.2) which are almost optimal, or minimal, for a spatial CLT. The result given also demonstrates the spatial block bootstrap for the sample mean under weaker mixing/moment conditions than considered previously (cf. Theorem 12.1, [19]).

5. Convolved subsampling in other contexts. We briefly outline relationships between convolved subsampling and some recent literature about block resampling for statistics outside of the sample mean cases in Sections 4.2–4.5. As alternatives to bootstrap, such works have considered generalized approaches to resampling that are convolved subsampling. When viewed as such, these previous

developments may be unified and simplified by general results here for mixing time series (or processes) (cf. Section 4.1), as illustrated in Section 5.1 for U-statistics and Section 5.2 for spectral estimators. Section 5.3 mentions extensions to further statistics, such as L-estimators.

For the classes of statistics next considered, our results with convolved subsampling cannot be used to directly justify the block bootstrap. However, our findings may still contribute to this end, as explained in Section 8.

5.1. *U-statistics.* U-statistics are a class of nonlinear functionals for prescribing statistics, such as the sample variance. Suppose that X_1, \dots, X_n arise from a stationary process and, based on a symmetric kernel $h : \mathbb{R}^2 \rightarrow \mathbb{R}$, define a (bivariate) U-statistic as

$$t_n \equiv t_n(X_1, \dots, X_n) = \frac{2}{n(n-1)} \sum_{1 \leq i < j \leq n} h(X_i, X_j),$$

which estimates a target parameter $t(P) \equiv \int \int h(x, y) dG(x) dG(y)$, where G denotes the marginal distribution of X_t . Consider the problem of estimating the distribution of $T_n \equiv \sqrt{n}(t_n - t(P))$, with scaling $\tau_n = \sqrt{n}$, under weak time dependence. The subsampling distribution $S_{n,\text{SUB}}$ is defined by computing the U-statistic $t_{n,b,i} = [b(b-1)]^{-1} 2 \sum_{i \leq j_1 < j_2 \leq i+b-1} h(X_{j_1}, X_{j_2})$ on each length b subsample $\{(X_i, \dots, X_{i+b-1})\}_{i=1}^{N_n \equiv n-b+1}$ in (2.1). In contrast, block bootstrap versions of U-statistics have a formulation similar to (2.5); see Dehling and Wendler [9], Sharipov and Wendler [33] and Leucht [23]. That is, a bootstrap sample $X_1^*, \dots, X_{n_1}^*$, $n_1 = k_n b$, is generated by resampling k_n blocks of length b (typically $k_n = \lfloor n/b \rfloor$) and then the U-statistic $t_{n_1}^* \equiv t_{n_1}(X_1^*, \dots, X_{n_1}^*)$ is calculated from the complete bootstrap sample to create a bootstrap rendition $T_n^* = \sqrt{n_1}(t_{n_1}^* - E_* t_{n_1}^*)$ of T_n . In this setting, the bootstrap distribution T_n^* would not generally correspond to that of a k_n -fold convolution C_{n,k_n} of the subsampling distribution $S_{n,\text{SUB}}$, as occurred in the sample mean case (Section 2.2).

However, Sharipov, Tewes and Wendler [32] recently considered an alternative block resampling estimator for U-statistics, which matches the convolved subsampling estimator C_{n,k_n} here based on the subsampling estimator $S_{n,\text{SUB}}$ for T_n described above. Note that, for stationary mixing data, Dehling and Wendler [9] (Theorem 1.8–Lemma 3.6) provide a CLT for the relevant U-statistic: $T_n \xrightarrow{d} N(0, \sigma^2)$ and $ET_n^2 \rightarrow \sigma^2$ as $n \rightarrow \infty$ where $\sigma^2 \equiv 4 \sum_{k=-\infty}^{\infty} \text{Cov}(h_1(X_0), h_1(X_k))$ for $h_1(x) = \int h(x, y) dG(y)$. Under mixing conditions and with $k_n = \lfloor n/b \rfloor \rightarrow \infty$, Sharipov, Tewes and Wendler [32] established that C_{n,k_n} captures this limiting normal distribution of T_n and also showed the consistency of the variance $\hat{\sigma}_{n,\text{SUB}}^2$ of C_{n,k_n} . The argument there involved decomposing the bootstrap U-statistic T_n^* into a linear part, coinciding with a sample mean from the usual block bootstrap, and degenerate part shown to be negligible. However, the general convolution result in Theorem 4 for mixing processes provides an alternative, and much simpler,

approach. From $T_n \xrightarrow{d} N(0, \sigma^2)$ and $ET_n^2 \rightarrow \sigma^2$, all of the conditions of Theorem 4 automatically hold, proving that C_{n,k_n} is consistent for the distribution of T_n for any convolution level k_n and also that $\hat{\sigma}_{n,\text{SUB}}^2 \xrightarrow{P} \sigma^2$. This approach also weakens the block assumptions used by [32] [i.e., $b = O(n^\epsilon)$ for some $\epsilon \in (0, 1)$] to $b^{-1} + b/n \rightarrow 0$ under Theorem 4.

5.2. *Spectral estimators for nonstationary time series.* As described in Section 4.2, almost periodically correlated (APC) time series $\{X_t\}$ are an important example of nonstationary sequences. Beyond the mean function, inference about the correlation structure is also of interest. Based on a sample X_1, \dots, X_n , a symmetric kernel $w(\cdot)$ and a bandwidth choice L_n , Lenart [21, 22] considered kernel estimators

$$t_n(X_1, \dots, X_n) \equiv \frac{1}{2\pi n} \sum_{t=1}^n \sum_{s=1}^n \frac{1}{L_n} w\left(\frac{t-s}{L_n}\right) X_t X_s e^{-i\nu t} e^{i\omega s}$$

for an extended spectral density $t(P) \equiv t(P)(\nu, \omega)$, $(\nu, \omega) \in (0, 2\pi]^2$, used to represent the almost periodic covariance function $c_\tau(t) \equiv \text{Cov}(X_t, X_{t+\tau})$, $t \in \mathbb{Z}$, for a given $\tau \in \mathbb{Z}$; see [21, 22] for details.

For $T_n \equiv \tau_n(t_n - t(P))$ with scaling $\tau_n = \sqrt{n/L_n}$, Lenart [21] (Theorems 3.1–3.2) proved a CLT $T_n \xrightarrow{d} N(0, \sigma^2)$ and moment convergence $ET_n^2 \rightarrow \sigma^2$ with mixing APC series, which was extended in Lenart [22] to multivariate data. Due to the complicated form of σ^2 , a subsampling estimator $S_{n,\text{SUB}}$ for the distribution of T_n may be computed as in (2.1) with analog statistics $t_{n,b,i}$ and scaling $\tau_b = \sqrt{b/L_b}$ defined from subsamples $\{(X_i, \dots, X_{i+b-1})\}_{i=1}^{N_n \equiv n-b+1}$. Lenart [21] proved the consistency of the estimator $S_{n,\text{SUB}}$, while Lenart [22] proposed a generalized resampling method which essentially corresponds to a convolved subsampling estimator C_{n,k_n} induced from $S_{n,\text{SUB}}$. In particular, Lenart [22] established the consistency of C_{n,k_n} through bootstrap arguments requiring much stronger mixing and moment assumptions than needed for the convergence $T_n \xrightarrow{d} N(0, \sigma^2)$ and $ET_n^2 \rightarrow \sigma^2$. However, the general convolved subsampling result in Theorem 4 may alternatively be used here with mixing nonstationary ACP series.

To apply Theorem 4 with blocks where $b^{-1} + b/n + \tau_b/\tau_n \rightarrow 0$ as $n \rightarrow \infty$, one requires that $Y_b \xrightarrow{d} N(0, \sigma^2)$ and that (3.2) holds, where Y_b , $b \equiv b_n \geq 1$, denotes a sequence of variables with distribution $D_{n,b}(\cdot)$ from (3.1). But, the same conditions needed for $T_n \xrightarrow{d} N(0, \sigma^2)$ and $ET_n^2 \rightarrow \sigma^2$ also yield $Y_b \xrightarrow{d} N(0, \sigma^2)$ and $EY_b^2 \rightarrow \sigma^2$ (cf. Theorems 3.1–3.2 and 4.1, [21]). Furthermore, mixing and $Y_b \xrightarrow{d} N(0, \sigma^2)$, along with $T_n = O_p(1)$ and $\tau_n/\tau_b \rightarrow \infty$, guarantee that (2.3) holds [i.e., $\sup_{x \in \mathbb{R}} |S_{n,\text{SUB}}(x) - \Phi(x/\sigma)| \xrightarrow{P} 0$] and that consequently (3.2) follows from Theorem 3(ii) by $EY_b^2 \rightarrow \sigma^2$. That is, the same minimal conditions for a CLT with APC series suffice for the consistency of convolved subsampling C_{n,k_n} by the general result of Theorem 4.

5.3. *L-estimators and other statistics beyond the sample mean.* Other classes of statistics with normal limits, where resampling may be helpful, include M-estimators, L-statistics and generalizations such as GL-statistics. Just as for the U-statistics and spectral estimators in the previous sections, convolved subsampling may be applied as a resampling method which is neither classical subsampling nor the block bootstrap for general statistics. As an advantage in such cases, convolved subsampling is often verifiable under mild assumptions (cf. Sections 3, 4.1), which we mention for L-estimators. If X_1, \dots, X_n denotes a stationary stretch with marginal quantile function G^{-1} , an L-estimator $t_n = \int_0^1 \hat{G}_n^{-1}(u)J(u) du$ of a parameter $t(P) = \int_0^1 G^{-1}(u)J(u) du$ represents a linear combinations of order statistics, defined by the quantiles of the empirical distribution $\hat{G}_n(x) = n^{-1} \sum_{t=1}^n I(X_t \leq x)$, $x \in \mathbb{R}$, and a weighting function $J : [0, 1] \rightarrow \mathbb{R}$. Under the mild mixing/moment conditions used for sample means in Theorem 5, the same conclusions for subsampling and convolved subsampling also hold with general L-estimators $T_n = \sqrt{n}(t_n - t(P))$, provided that J is bounded and continuous almost everywhere; see the Supplementary Material [15] for more formal details regarding L-estimators. Section 7 provides some numerical justification of the convolved procedure for the trimmed mean as an L-statistic.

6. Independent data versions. For completeness, we briefly mention a variation of convolved subsampling for independent data. Recall that Section 4.1 considered block-based convolved subsampling with general statistics computed from strongly mixing time processes $\{X_t\}$. Hence, results of Section 4.1 apply to independent data, as do block bootstrap results of Section 4.2 for sample means under mixing conditions. However, with independent X_1, \dots, X_n , one may consider a different formulation of subsamples rather than data blocks of b consecutive observations. Namely, let $b_n \equiv b$ denote a set size and define subsamples $Y_{b,1}, \dots, Y_{b,N_n}$ as the $N_n \equiv \binom{n}{b}$ unordered subsets of size b from $\{X_1, \dots, X_n\}$. The “independent data” subsampling estimator $S_{n,\text{SUB}}^{\text{ID}}$ is defined as $S_{n,\text{SUB}}$ in (2.1) with subsample statistics $t_{n,b,i} \equiv t_b(Y_{b,i})$, $i = 1, \dots, N_n$, where statistics $t_b(\cdot)$ are symmetric in their arguments here; see Politis, Romano and Wolf [29] (Chapter 2) for a general treatment of this subsampling estimator with i.i.d. data.

The next theorem verifies that, for independent data, the general results for convolved subsampling with previous block-based subsamples (Section 4.1) also hold when the convolution is based on the independent data subsampling estimator $S_{n,\text{SUB}}^{\text{ID}}$ using all subsets of size b .

THEOREM 6. *Let $\{X_t\}$ be a sequence of independent (possibly non-i.i.d.) random variables. Given $S_{n,\text{SUB}}^{\text{ID}}$, let $(\hat{\sigma}_{n,\text{SUB}}^{\text{ID}})^2$ and C_{n,k_n}^{ID} denote the corresponding subsampling variance estimator and convolved subsampling estimator. Then Theorem 4 holds under the notational convention that $S_{n,\text{SUB}} \equiv S_{n,\text{SUB}}^{\text{ID}}$, $\hat{\sigma}_{n,\text{SUB}}^2 \equiv$*

$(\hat{\sigma}_{n,\text{SUB}}^{\text{ID}})^2$, and $C_{n,k_n} \equiv C_{n,k_n}^{\text{ID}}$ and that subsample quantities in (3.1)–(3.2) are defined as $T_{b,i} = \tau_b[t_b(Y_{b,i}) - t(P)]$, $i = 1, \dots, N_n \equiv \binom{n}{b}$.

Additionally, if the variables $\{X_i\}$ are i.i.d., then Corollary 2 likewise holds.

We may also draw some connections between convolved subsampling and the bootstrap for sample means with independent data. Suppose independent variables X_1, \dots, X_n have common mean μ (e.g., as in Tukey’s symmetric contamination model where observations may have different variances), from which we define $T_n \equiv \sqrt{n}(\bar{X}_n - \mu)$. The convolved estimator C_{n,k_n}^{ID} here has close parallels to the classic independent bootstrap of Efron [11]. Namely, C_{n,k_n}^{ID} is the resampling distribution of $T_n^* \equiv \sqrt{n_1}(\bar{X}_{n_1}^* - \bar{X}_n)$ for a sample mean $\bar{X}_{n_1}^*$ of size $n_1 = k_n b$ formed by averaging k_n independent subsamples of size b , with each size b subsample drawn uniformly and without replacement from $\{X_i\}_{i=1}^n$; if the subsamples of size b are instead drawn with replacement from $\{X_i\}_{i=1}^n$, then T_n^* alternatively produces the independent bootstrap distribution, say $C_{n,k_n b}^{\text{ID,boot}}$, with a resample size $n_1 = k_n b$. Consequently, the independent data version of convolved subsampling C_{n,k_n}^{ID} does not exactly match the independent bootstrap. However, the following result for independent data shows that the subsampling estimator $S_{n,\text{SUB}}^{\text{ID}}$ and its convolution C_{n,k_n}^{ID} are valid in a broad non-i.i.d. context for sample means, and the differences between C_{n,k_n}^{ID} and $C_{n,k_n b}^{\text{ID,boot}}$ are asymptotically negligible.

THEOREM 7. *Let X_1, X_2, \dots , denote a sequence of independent (possibly non-i.i.d.) variables, with finite variances and common mean $\text{EX}_t = \mu \in \mathbb{R}$. Define $X_{i,\mu} \equiv X_i - \mu$, $i \geq 1$. As $n \rightarrow \infty$, suppose $b^{-1} + b/n \rightarrow 0$ and that*

$$\frac{1}{n} \sum_{i=1}^n \text{EX}_{i,\mu}^2 I[|X_{i,\mu}| > \epsilon\sqrt{b}] \rightarrow 0$$

and

$$\max_{1 \leq i_1 < i_2 < \dots < i_b \leq n} \left| \frac{1}{b} \sum_{j=1}^b \text{EX}_{i_j,\mu}^2 - \sigma^2 \right| \rightarrow 0$$

for each $\epsilon > 0$ and some $\sigma^2 > 0$. Then, as $n \rightarrow \infty$, $T_n = \sqrt{n}(\bar{X}_n - \mu) \xrightarrow{d} N(0, \sigma^2)$ along with $\sup_{x \in \mathbb{R}} |S_{n,\text{SUB}}^{\text{ID}}(x) - \Phi(x/\sigma)| \xrightarrow{p} 0$ and $(\hat{\sigma}_{n,\text{SUB}}^{\text{ID}})^2 \xrightarrow{p} \sigma^2$. Furthermore, for any positive integer sequence k_n ,

$$\sup_{x \in \mathbb{R}} |C_{n,k_n}^{\text{ID}}(x) - \Phi(x/\sigma)| \xrightarrow{p} 0 \quad \text{and} \quad d_2[C_{n,k_n}^{\text{ID}}, C_{n,k_n b}^{\text{ID,boot}}] \xrightarrow{p} 0,$$

where $d_2(\cdot, \cdot)$ denotes Mallow’s metric between distributions C_{n,k_n}^{ID} and $C_{n,k_n b}^{\text{ID,boot}}$.

REMARK 4. Above $d_2(\cdot, \cdot)$ metricizes weak convergence [5], where, for distributions F and G on \mathbb{R} , $[d_2(F, G)]^2 \equiv \inf\{E|X - Y|^2 : X \sim F, Y \sim G\}$ with the infimum over all pairs (X, Y) with marginal distributions F and G .

A Lindeberg condition, defined by replacing b with n in the second moment assumptions of Theorem 7, suffices for $T_n = \sqrt{n}(\bar{X}_n - \mu) \xrightarrow{d} N(0, \sigma^2)$. Hence, Theorem 7 validates subsampling, convolved subsampling and the bootstrap for the sample mean under a slightly stronger condition than required for the CLT with independent data. This finding also involves a weaker moment condition than a classical bootstrap result of Liu [24] for sample means of non-i.i.d. data (i.e., $\sup_{i \geq 1} E|X_i|^{2+\delta} < \infty$ with $\delta > 0$). Also, for general statistics with i.i.d. data, Politis, Romano and Wolf [29] (Corollary 2.3.1) use the subsampling estimator $S_{n, \text{SUB}}^{\text{ID}}$ to prove the consistency of a version of the bootstrap with a resample size of $b < n$, provided that $b^{-1} + b^2/n \rightarrow 0$. When specialized to sample means, their intended bootstrap becomes $C_{n, k_n b}^{\text{ID, boot}}$ with convolution $k_n = 1$ and Theorem 7 generalizes their result under a weaker requirement $b/n \rightarrow 0$ and for non-i.i.d. data.

7. Numerical results. Particularly in the important case of the sample mean, where convolved subsampling equals the classical block bootstrap, many authors have shown, via numerical as well as theoretical evidence, that the bootstrap is generally advantageous over standard subsampling when both are valid (cf. [19], Chapter 6; [29], Chapter 10). Intuitively, the i.i.d. randomization of the bootstrap often better aligns its distribution with a normal target. Hence, when considering the sample mean, our aim here is of theoretical nature and we provide a new approach for proving bootstrap consistency.

For statistics of interest beyond the sample mean, the situation is different. In this scenario, convolved subsampling estimation gives a completely new resampling procedure, where results from Section 3 can be used to establish the method’s consistency in general contexts. As described in Section 5, the method has been applied to spectral densities [21] and U-statistics [32], where some numerical findings have indicated improvements over subsampling with U-statistics. Outside of this, not much is presently known about the method’s properties relative to subsampling or block bootstrap. However, for approximating normal limits, convolved subsampling might generally be anticipated to perform better than subsampling, as convolution may move estimators closer to normality (cf. Remark 5 to follow).

For comparison, we include a small simulation study involving the trimmed mean statistic (cf. Section 5.3), computed from a time series X_1, \dots, X_n as

$$\bar{X}_{n, \tau_1, \tau_2} = \frac{1}{[n\tau_2] - [n\tau_1]} \sum_{i=[n\tau_1]+1}^{[n\tau_2]} X_{(i)},$$

using order statistics $X_{(1)}, \dots, X_{(n)}$ and trimming proportions $0 \leq \tau_1 < \tau_2 \leq 1$. When $\tau_1 = 0$ and $\tau_2 = 1$, the trimmed mean equals the sample mean and, conse-

quently, convolved subsampling becomes equivalent to the block bootstrap. However, when observations are discarded by $\tau_1 \neq 0$ or $\tau_2 \neq 1$, the two resampling procedures are different. For the trimmed mean parameter, 90% confidence intervals are given by $\bar{X}_{n,\tau_1,\tau_2} \pm q_{0.9}/\sqrt{n}$, with $q_{0.9}$ denoting an appropriate quantile found by either subsampling, convolved subsampling with different levels of convolution, or the moving blocks bootstrap. For data generation, we use an AR(1)-process with standard normal marginals.

Table 1 displays the coverage probabilities from intervals by the aforementioned resampling methods, based on an AR(1) parameter 0.3 and various sample sizes n ;

TABLE 1
 Coverage percentages of 90% intervals for the trimmed mean based on subsampling ($S_{n,SUB}$), convolved subsampling (C_{n,k_n}) and block bootstrap ($C_{n,\lfloor n/b \rfloor}^{boot}$) for various block sizes b , convolution levels k_n and sample sizes n ; coverages are based on 1000 simulations from an AR(1)-process (coefficient 0.3) with distributions for C_{n,k_n} or $C_{n,\lfloor n/b \rfloor}^{boot}$ approximated from 999 resamples. Note $C_{n,\lfloor n/b \rfloor}$ and $C_{n,\lfloor n/b \rfloor}^{boot}$ match for $\tau_1 = 1, \tau_2 = 0$

n	50		100		200		500		1000	
	7	3	10	4	14	5	22	7	31	9
Percentages of trimming $\tau_1 = 1 - \tau_2 = 0$										
$S_{n,SUB}$	79.1	84.5	84.2	86.3	85.0	86.9	89.3	88.9	87.3	89.1
$C_{n,3}$	80.1	84.8	85.3	86.5	84.2	87.2	89.4	88.5	87.3	88.8
$C_{n,\lfloor n/(2b) \rfloor}$	80.3	85.0	86.1	86.7	85.3	87.5	89.4	89.1	87.7	89.3
$C_{n,\lfloor n/b \rfloor}$	82.0	85.5	86.4	87.4	85.5	87.8	89.4	89.2	87.6	89.9
$C_{n,\lfloor n/b \rfloor}^{boot}$	82.0	85.5	86.4	87.4	85.5	87.8	89.4	89.2	87.6	89.9
Percentages of trimming $\tau_1 = 1 - \tau_2 = 0.05$										
$S_{n,SUB}$	82.4	86.3	83.9	87.6	84.3	88.4	88.0	87.1	87.3	87.3
$C_{n,3}$	82.3	85.9	84.4	88.4	83.7	89.1	88.0	86.7	87.4	87.8
$C_{n,\lfloor n/(2b) \rfloor}$	82.5	86.2	84.7	87.9	84.2	89.2	88.0	87.7	87.4	88.0
$C_{n,\lfloor n/b \rfloor}$	84.2	87.4	85.1	88.7	84.3	89.8	88.3	88.1	87.6	87.7
$C_{n,\lfloor n/b \rfloor}^{boot}$	87.2	81.3	84.9	85.7	86.0	85.7	89.4	89.6	88.4	88.9
Percentages of trimming $\tau_1 = 1 - \tau_2 = 0.10$										
$S_{n,SUB}$	80.5	85.3	83.8	85.2	84.9	86.4	87.2	86.7	88.5	89.1
$C_{n,3}$	80.5	86.4	83.8	85.4	84.6	86.9	87.1	87.1	88.7	88.9
$C_{n,\lfloor n/(2b) \rfloor}$	80.7	86.5	84.5	85.6	84.9	87.7	86.8	87.0	88.3	89.2
$C_{n,\lfloor n/b \rfloor}$	82.2	87.3	85.4	85.9	86.1	88.0	88.0	87.9	88.2	89.4
$C_{n,\lfloor n/b \rfloor}^{boot}$	82.4	82.1	85.3	84.8	84.7	86.7	88.8	86.4	89.5	87.8
Percentages of trimming $\tau_1 = 1 - \tau_2 = 0.25$										
$S_{n,SUB}$	80.3	84.3	85.0	85.9	84.1	84.1	86.2	85.2	85.8	88.0
$C_{n,3}$	80.7	85.1	85.7	86.2	84.5	84.2	86.5	85.0	86.1	87.0
$C_{n,\lfloor n/(2b) \rfloor}$	81.7	86.0	85.8	86.6	85.4	85.1	86.3	85.3	86.3	87.7
$C_{n,\lfloor n/b \rfloor}$	82.8	86.3	86.8	87.8	85.9	85.9	87.0	85.9	86.5	88.9
$C_{n,\lfloor n/b \rfloor}^{boot}$	82.6	81.8	84.8	86.7	87.1	87.6	88.6	86.7	88.9	89.9

TABLE 2
Lengths of 90% intervals for the trimmed mean corresponding to the coverage probabilities reported in Table 1

<i>n</i>	50		100		200		500		1000	
	7	3	10	4	14	5	22	7	31	9
Percentages of trimming $\tau_1 = 1 - \tau_2 = 0$										
$S_{n,SUB}$	0.570	0.606	0.423	0.435	0.309	0.315	0.202	0.202	0.144	0.144
$C_{n,3}$	0.598	0.635	0.438	0.452	0.314	0.322	0.205	0.206	0.145	0.146
$C_{n, \lfloor n/(2b) \rfloor}$	0.580	0.623	0.430	0.445	0.310	0.318	0.203	0.204	0.145	0.145
$C_{n, \lfloor n/b \rfloor}$	0.578	0.615	0.425	0.440	0.306	0.315	0.202	0.202	0.144	0.144
$C_{n, \lfloor n/b \rfloor}^{boot}$	0.578	0.615	0.425	0.440	0.306	0.315	0.202	0.202	0.144	0.144
Percentages of trimming $\tau_1 = 1 - \tau_2 = 0.05$										
$S_{n,SUB}$	0.567	0.605	0.430	0.436	0.310	0.315	0.202	0.202	0.144	0.145
$C_{n,3}$	0.595	0.633	0.444	0.450	0.316	0.321	0.204	0.205	0.146	0.146
$C_{n, \lfloor n/(2b) \rfloor}$	0.576	0.623	0.437	0.444	0.312	0.317	0.203	0.203	0.145	0.145
$C_{n, \lfloor n/b \rfloor}$	0.575	0.614	0.432	0.439	0.310	0.315	0.201	0.203	0.144	0.145
$C_{n, \lfloor n/b \rfloor}^{boot}$	0.594	0.580	0.446	0.425	0.317	0.309	0.205	0.200	0.146	0.144
Percentages of trimming $\tau_1 = 1 - \tau_2 = 0.10$										
$S_{n,SUB}$	0.571	0.605	0.423	0.432	0.313	0.314	0.202	0.202	0.147	0.144
$C_{n,3}$	0.597	0.636	0.437	0.450	0.320	0.321	0.205	0.204	0.148	0.146
$C_{n, \lfloor n/(2b) \rfloor}$	0.577	0.624	0.428	0.443	0.315	0.318	0.203	0.203	0.147	0.145
$C_{n, \lfloor n/b \rfloor}$	0.578	0.615	0.425	0.437	0.312	0.314	0.202	0.202	0.147	0.144
$C_{n, \lfloor n/b \rfloor}^{boot}$	0.603	0.586	0.442	0.430	0.322	0.311	0.206	0.202	0.149	0.145
Percentages of trimming $\tau_1 = 1 - \tau_2 = 0.25$										
$S_{n,SUB}$	0.571	0.607	0.432	0.437	0.323	0.315	0.209	0.203	0.151	0.148
$C_{n,3}$	0.602	0.647	0.453	0.460	0.333	0.329	0.214	0.210	0.153	0.152
$C_{n, \lfloor n/(2b) \rfloor}$	0.578	0.627	0.440	0.448	0.326	0.322	0.211	0.206	0.151	0.150
$C_{n, \lfloor n/b \rfloor}$	0.578	0.614	0.434	0.439	0.323	0.316	0.209	0.203	0.150	0.148
$C_{n, \lfloor n/b \rfloor}^{boot}$	0.623	0.614	0.459	0.451	0.335	0.326	0.214	0.211	0.153	0.151

qualitatively similar results with other AR parameters appear in the Supplementary Material [15]. Additionally, corresponding interval lengths are reported in Table 2. As those differ only marginally over the five resampling methods, we will focus on the probabilities. In general, all methods tend to yield coverage probabilities below the nominal level of 90%. Moreover, convolution of subsampling improves performance over regular subsampling in almost all cases, with increasing levels of convolution tending to produce better coverage accuracy. For illustration, the subsampling distributional approximation for the trimmed mean statistic is shown in Figure 1 along with counterparts improved by convolution. In terms of comparisons between the block bootstrap and convolved subsampling, the methods again differ when $\tau_1 \neq 0$ or $\tau_2 \neq 1$, as indicated in Table 1. Based on our simulation, neither method seems to be clearly advantageous. Depending on the choice of

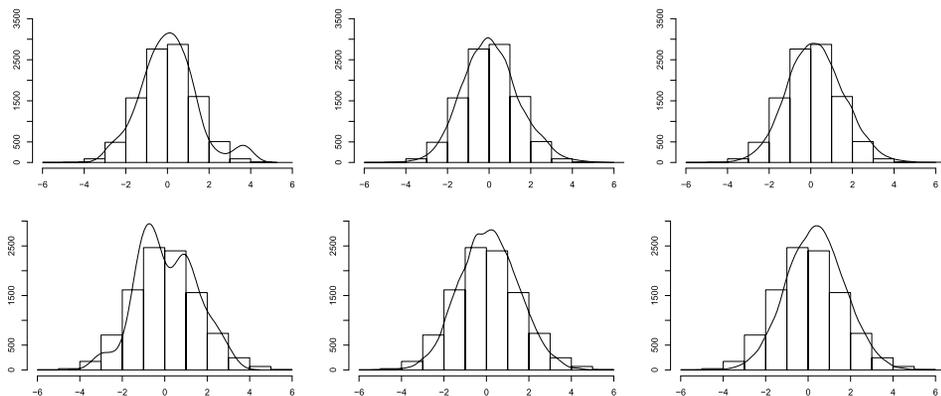


FIG. 1. Histogram of the centered/scaled trimmed mean, with $\tau_1 = 1 - \tau_2 = 0.10$ (top) or $\tau_1 = 1 - \tau_2 = 0.25$ (bottom), based on 10,000 simulations from an AR(1) process, and density estimates by convolved subsampling with convolution levels of $k_n = 1$ (i.e., subsampling) (left), $k_n = 3$ (center) and $k_n = 14$ (right) from one data realization.

trimming parameters, block length b , and level of convolution k_n , either bootstrap or convolved subsampling may emerge as the better of the two.

REMARK 5. For approximating normal limits, convolved subsampling may reduce skewness in distributional estimates from basic subsampling. For illustration, considering the sample mean, the distribution of $T_n = \sqrt{n}(\bar{X}_n - \mu)$ often has approximate skewness γ/\sqrt{n} for some constant γ . In this case, the corresponding subsampling estimator $S_{n,\text{SUB}}$ is known to have a larger approximate skewness γ/\sqrt{b} , while a more fully convolved estimator (bootstrap) C_{n,k_n} with $k_n \approx n/b$ has skewness approximately γ/\sqrt{n} . A better matching skewness may improve higher-order accuracy; see [29] (Section 10.2).

8. Concluding remarks and extensions. For approximating sampling distributions with normal limits, we have developed a theory for the k -fold self-convolution of subsampling estimators. Results validate the method for general statistics and dependent data structures, based on mild consistency properties of the basic subsampling estimator and its subsampling variance. The latter estimator is crucial under diverging levels of convolution, as occurs with block bootstrap. For time series, convolved subsampling matches the block bootstrap for sample means. With more general statistics, convolved subsampling often differs from the bootstrap and, instead, provides a hybrid-type of resampling that has received recent consideration (cf. Section 5) and may improve upon standard subsampling for normal targets. Further study is required of higher-order accuracy. However, as convolved subsampling can often be verified under mild process assumptions, this offers an alternative approach for establishing the block bootstrap under weaker conditions than previously considered for sample means in particular.

We make two final points about applying convolved subsampling to validate the block bootstrap with more general statistics, which is difficult to characterize. First, from the standard linearization technique based on functional representations and differentials, a target statistic may often be decomposed into main/linear and remainder parts (cf. [17]; Chapter 4.3, [19]). Commonly, the linear term is a sample mean (e.g., of influence functions), while the remainder is negligible. Convolved subsampling can be directly used to establish the block bootstrap for this main term under weak conditions. Our framework, though, does not immediately address the remainder.

However, second, such remainders are often smaller-order than a norm of the centered empirical distribution, in both the original data and bootstrap worlds, where the bootstrap version is, say, $\sqrt{n}\|\hat{G}_n^* - E_*\hat{G}_n^*\|$. By virtue of the empirical distribution as an average, the bootstrap estimator $\sqrt{n}(\hat{G}_n^*(x) - E_*\hat{G}_n^*(x))$, $x \in \mathbb{R}$, must consequently match convolved subsampling. This connection could potentially foster bootstrap studies with empirical processes and dependent data, particularly as subsampling estimators are consistent for empirical processes under weak assumptions ([29], Chapter 7.4). Further investigation is required, though, because the technical obstacle to weak convergence differs substantially from the usual sample mean case: namely, tightness of the bootstrap/convolved subsampling process $\sqrt{n}(\hat{G}_n^*(\cdot) - E_*\hat{G}_n^*(\cdot))$ must be determined on a suitable metric space (cf. [19], Chapter 4.4.1). Still, convolved subsampling may provide a general tool for advancing future developments with resampling dependent data.

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SUPPLEMENTARY MATERIAL

Supplement to “Convolved subsampling estimation with applications to block bootstrap” (DOI: [10.1214/18-AOS1695SUPP](https://doi.org/10.1214/18-AOS1695SUPP); .pdf). This supplement provides proofs for the distributional results about convolved subsampling and further numerical/theoretical support for the simulation study.

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