

Index sequences - 770220-96: Plate C (193-288)

When sequencing, assign 17 cycles for Index1 and 8 cycles for Index2.

Index1 (i7) Number	Index1 (i7) Sequence	Index1 (i7) Sequence + UMI	Index2 (i5) Number	Index2 (i5) Forward Sequence	Index2 (i5) Reverse Complement Sequence
193	CTTCGTTC	CTTCGTTCNNNNNNNNNN	693	ACCTTCTC	GAGAAGGT
194	GTCTAGGT	GTCTAGGTNNNNNNNNNN	694	TCGTGGAT	ATCCACGA
195	ACGTCGTA	ACGTCGTANNNNNNNNNN	695	GTTTCATGG	CCATGAAC
196	GAGCTCAA	GAGCTCAANNNNNNNNNN	696	TAGGATGC	GCATCCTA
197	CGTGTACT	CGTGTACTNNNNNNNNNN	697	CATGGAAC	GTTCCATG
198	CACTGACA	CACTGACANNNNNNNNNN	698	GCTTAGCT	AGCTAAGC
199	TCGTAGTC	TCGTAGTCNNNNNNNNNN	699	CTAACTCG	CGAGTTAG
200	GCACGTAA	GCACGTAANNNNNNNNNN	700	ACCATGTG	CACATGGT
201	CAAGCAGT	CAAGCAGTNNNNNNNNNN	701	TCAGACGA	TCGTCTGA
202	ACATAGGC	ACATAGGCNNNNNNNNNN	702	TATCAGCG	CGCTGATA
203	TGTGGTAC	TGTGGTACNNNNNNNNNN	703	AGCAGATG	CATCTGCT
204	CACCACTA	CACCACTANNNNNNNNNN	704	AACGGTCA	TGACCGTT
205	CTGCGTAT	CTGCGTATNNNNNNNNNN	705	CGAACTGT	ACAGTTCG
206	ACGGTCTT	ACGGTCTTNNNNNNNNNN	706	TCCGAGTT	AACTCGGA
207	GATTGGAG	GATTGGAGNNNNNNNNNN	707	TTCTCTCG	CGAGAGAA
208	TGTCCAGA	TGTCCAGANNNNNNNNNN	708	ATTCTGGC	GCCAGAAT
209	CCAGTGTT	CCAGTGTTNNNNNNNNNN	709	ACTGCTAG	CTAGCAGT

210	TGCACCAA	TGCACCAANNNNNNNNN	710	CATAACGG	CCGTTATG
211	TTGACAGG	TTGACAGGNNNNNNNNN	711	CAGTCTTC	GAAGACTG
212	AGGCATAG	AGGCATAGNNNNNNNNN	712	TGCCTCTT	AAGAGGCA
213	TAGCCGAA	TAGCCGAANNNNNNNNN	713	ACTGTGTC	GACACAGT
214	TTGTCGGT	TTGTCGGTNNNNNNNNN	714	GTATTGGC	GCCAATAC
215	CATCTACG	CATCTACGNNNNNNNNN	715	CGATGCTT	AAGCATCG
216	GCATACAG	GCATACAGNNNNNNNNN	716	AAGGCTGA	TCAGCCTT
217	ACAGCAAC	ACAGCAACNNNNNNNNN	717	AGTCAGGA	TCCTGACT
218	CTGGTTCT	CTGGTTCTNNNNNNNNN	718	CAGGTATC	GATACCTG
219	TCGACATC	TCGACATCNNNNNNNNN	719	TCTCCGAT	ATCGGAGA
220	AACCTCCT	AACCTCCTNNNNNNNNN	720	TTCAGCCT	AGGCTGAA
221	CAGCGATT	CAGCGATTNNNNNNNNN	721	TCTGAGAG	CTCTCAGA
222	AGGTCACT	AGGTCACTNNNNNNNNN	722	TTAGGTCG	CGACCTAA
223	GCAATTCG	GCAATTCGNNNNNNNNN	723	CTCTGGTT	AACCAGAG
224	GCTTCTTG	GCTTCTTGNNNNNNNNN	724	GCGTTCTA	TAGAACGC
225	AACTGGTG	AACTGGTGNNNNNNNNN	725	TCACGTTT	GAACGTGA
226	CGGAATAC	CGGAATACNNNNNNNNN	726	AGGATGGT	ACCATCCT
227	GCTTCGAA	GCTTCGAANNNNNNNNN	727	GTGTTCTT	AGGAACAC
228	CAAGGTCT	CAAGGTCTNNNNNNNNN	728	GTAGCATC	GATGCTAC
229	AACCTTGG	AACCTTGGNNNNNNNNN	729	AGGATCTG	CAGATCCT
230	CCATACGT	CCATACGTNNNNNNNNN	730	GACAAGAG	CTCTTGTC
231	TGGTCCTT	TGGTCCTTNNNNNNNNN	731	TTACGGCT	AGCCGTAA
232	ACCGCATA	ACCGCATANNNNNNNNN	732	GCTGTTGT	ACAACAGC
233	CCTTCCTT	CCTTCCTTNNNNNNNNN	733	AACCGAAG	CTTCGGTT
234	TACACGCT	TACACGCTNNNNNNNNN	734	TCTGCTCT	AGAGCAGA

235	TGCGTAGA	TGCGTAGANNNNNNNNN	735	CTCAGCTA	TAGCTGAG
236	AAGAGCCA	AAGAGCCANNNNNNNNN	736	CTTCACCA	TGGTGAAG
237	ATGGAAGG	ATGGAAGGNNNNNNNNN	737	GATCGTAC	GTACGATC
238	GCCAGTAT	GCCAGTATNNNNNNNNN	738	CTACAGTG	CACTGTAG
239	CGTAGGTT	CGTAGGTTNNNNNNNNN	739	TCGAGTGA	TCACTCGA
240	CGAGTATG	CGAGTATGNNNNNNNNN	740	CAAGTGCA	TGCACTTG
241	CAAGTGCA	CAAGTGCANNNNNNNNN	741	CGAGTATG	CATACTCG
242	TCGAGTGA	TCGAGTGANNNNNNNNN	742	CGTAGGTT	AACCTACG
243	CTACAGTG	CTACAGTGNNNNNNNNN	743	GCCAGTAT	ATACTGGC
244	GATCGTAC	GATCGTACNNNNNNNNN	744	ATGGAAGG	CCTTCCAT
245	CTTCACCA	CTTCACCANNNNNNNNN	745	AAGAGCCA	TGGCTCTT
246	CTCAGCTA	CTCAGCTANNNNNNNNN	746	TGCGTAGA	TCTACGCA
247	TCTGCTCT	TCTGCTCTNNNNNNNNN	747	TACACGCT	AGCGTGTA
248	AACCGAAG	AACCGAAGNNNNNNNNN	748	CCTTCCTT	AAGGAAGG
249	GCTGTTGT	GCTGTTGTNNNNNNNNN	749	ACCGCATA	TATGCGGT
250	TTACGGCT	TTACGGCTNNNNNNNNN	750	TGGTCCTT	AAGGACCA
251	GACAAGAG	GACAAGAGNNNNNNNNN	751	CCATACGT	ACGTATGG
252	AGGATCTG	AGGATCTGNNNNNNNNN	752	AACCTTGG	CCAAGGTT
253	GTAGCATC	GTAGCATCNNNNNNNNN	753	CAAGGTCT	AGACCTTG
254	GTGTTCCCT	GTGTTCCCTNNNNNNNNN	754	GCTTCGAA	TTCGAAGC
255	AGGATGGT	AGGATGGTNNNNNNNNN	755	CGGAATAC	GTATTCCG
256	TCACGTTC	TCACGTTCNNNNNNNNN	756	AACTGGTG	CACCAGTT
257	GCGTTCTA	GCGTTCTANNNNNNNNN	757	GCTTCTTG	CAAGAAGC
258	CTCTGGTT	CTCTGGTTNNNNNNNNN	758	GCAATTCG	CGAATTGC
259	TTAGGTCG	TTAGGTCGNNNNNNNNN	759	AGGTCACT	AGTGACCT

260	TCTGAGAG	TCTGAGAGNNNNNNNNNN	760	CAGCGATT	AATCGCTG
261	TTCAGCCT	TTCAGCCTNNNNNNNNNN	761	AACCTCCT	AGGAGGTT
262	TCTCCGAT	TCTCCGATNNNNNNNNNN	762	TCGACATC	GATGTCGA
263	CAGGTATC	CAGGTATCNNNNNNNNNN	763	CTGGTTCT	AGAACCAG
264	AGTCAGGA	AGTCAGGANNNNNNNNNN	764	ACAGCAAC	GTTGCTGT
265	AAGGCTGA	AAGGCTGANNNNNNNNNN	765	GCATACAG	CTGTATGC
266	CGATGCTT	CGATGCTTNNNNNNNNNN	766	CATCTACG	CGTAGATG
267	GTATTGGC	GTATTGGCNNNNNNNNNN	767	TTGTCCGT	ACCGACAA
268	ACTGTGTC	ACTGTGTCNNNNNNNNNN	768	TAGCCGAA	TTCGGCTA
269	TGCCTCTT	TGCCTCTTNNNNNNNNNN	769	AGGCATAG	CTATGCCT
270	CAGTCTTC	CAGTCTTCNNNNNNNNNN	770	TTGACAGG	CCTGTCAA
271	CATAACGG	CATAACGGNNNNNNNNNN	771	TGCACCAA	TTGGTGCA
272	ACTGCTAG	ACTGCTAGNNNNNNNNNN	772	CCAGTGTT	AACACTGG
273	ATTCTGGC	ATTCTGGCNNNNNNNNNN	773	TGTCCAGA	TCTGGACA
274	TTCTCTCG	TTCTCTCGNNNNNNNNNN	774	GATTGGAG	CTCCAATC
275	TCCGAGTT	TCCGAGTTNNNNNNNNNN	775	ACGGTCTT	AAGACCGT
276	CGAACTGT	CGAACTGTNNNNNNNNNN	776	CTGCGTAT	ATACGCAG
277	AACGGTCA	AACGGTCANNNNNNNNNN	777	CACCACTA	TAGTGGTG
278	AGCAGATG	AGCAGATGNNNNNNNNNN	778	TGTGGTAC	GTACCACA
279	TATCAGCG	TATCAGCGNNNNNNNNNN	779	ACATAGGC	GCCTATGT
280	TCAGACGA	TCAGACGANNNNNNNNNN	780	CAAGCAGT	ACTGCTTG
281	ACCATGTG	ACCATGTGNNNNNNNNNN	781	GCACGTAA	TTACGTGC
282	CTAACTCG	CTAACTCGNNNNNNNNNN	782	TCGTAGTC	GACTACGA
283	GCTTAGCT	GCTTAGCTNNNNNNNNNN	783	CACTGACA	TGTCAGTG
284	CATGGAAC	CATGGAACNNNNNNNNNN	784	CGTGTACT	AGTACACG

285	TAGGATGC	TAGGATGCNNNNNNNNNN	785	GAGCTCAA	TTGAGCTC
286	GTTTCATGG	GTTTCATGGNNNNNNNNNN	786	ACGTCGTA	TACGACGT
287	TCGTGGAT	TCGTGGATNNNNNNNNNN	787	GTCTAGGT	ACCTAGAC
288	ACCTTCTC	ACCTTCTCNNNNNNNNNN	788	CTTCGTTC	GAACGAAG

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