Supplemental Methods: Primers used in this study. (The specific multiplex tag sequences used are listed in the GEO data submission file).

UPTAG Primer Sequences

(Illumina Adaptor sequence, unique multiplex tag, U1):
AATGATACGGCGACCACCGAGATCTACACTCTTTCCCTACACGACGCTCTTCCGATCTxxx
GATGTCCACGAGGTCTCT

U1 UPTAG primers above were used with a modified U2c* kanMX sequence:
CAAGCAGAAGACGGCATACGAGCTCTTCCGATCTtggatgcaaatgattatacatgg
(Illumina Adaptor sequence, kanMX sequence) to amplify the YKO UPTAG barcodes from each sorted population. The size of the amplicons from this PCR reaction is ~350bp.

DNTAG Primer Sequences

(Illumina Adaptor sequence, unique multiplex tag, D1)

AATGATACGGCGACCACCGAGATCTACACTCTTTCCCTACACGACGCTCTTCCGATCTxxx
CGGTGTCGGTCTCGTAG

D1 DNTAG primers above were amplified with a modified D2c* sequence:
CAAGCAGAAGACGGCATACGAGCTCTTCCGATCTtaacgccgccatccagtgtcg (Illumina Adaptor sequence, *kanMX* sequence) to amplify the YKO DNTAG barcodes from each sorted population. The size of the amplicon from this PCR reaction is ~180bp.

References

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