

Table S3. nCounter® Human Reference GX kit probe set.

Gene	Accession Number	Sequence
ABCF1	NM_001090.2	GATGTCCTCCCGCAAGCCATGTTAGAAAAATGCATCTGACATCAAGCTGGAGAAGTTCAGCATCTCCGCTCATGGCAAGGAGCTGTTTCGTCAATGCAGAC
ACTB	NM_001101.2	TGCAGAAGGAGATCACTGCCCTGGCACCCAGCACAAATGAAGATCAAGATCATTGCTCCTCCTGAGCGCAAGTACTCCGTGTGGATCGGCGGCTCCATCCT
ALAS1	NM_000688.4	GGGGATCGGGATGGAGTCATGCCAAAATGGACATCATTCTGGAACTTGGCAAAGCCTTTGGTTGTGTTGGAGGGTACATGCCAGCAGCAGTTCTC
B2M	NM_004048.2	CGGGCATTCTGAAGCTGACAGCATTGCGGCCGAGATGTCTCGCTCCGTGGCCTTAGCTGTGCTCGCGCTACTCTCTCTTTCTGGCCTGGAGGCTATCCA
CLTC	NM_004859.2	GGGTATCAACCCAGCAAAACATTGGCTTCAGTACCCTGACTATGGAGTCTGACAAATTCATCTGCATTAGAGAAAAAGTAGGAGAGCAGGCCAGGTGGTA
G6PD	NM_000402.2	ACAACATCGCCTGCGTTATCCTCACCTTCAAGGAGCCCTTTGGCACTGAGGGTCCGCGGGGGCTATTTGCATGAATTTGGGATCATCCGGACGTGATGCA
GAPDH	NM_002046.3	TCCTCCTGTTTCGACAGTCAGCCGCATCTCTTTTGCCTGCCAGCCGAGCCACATCGCTCAGACACCATGGGAAGGTGAAGGTCGGAGTCAACCGATTT
GUSB	NM_000181.1	CGGTCGTGATGTGGTCTGTGGCCAAAGAGCCTGCGTCCACCTAGAATCTGCTGGCTACTACTTGAAGATGGTATCGCTCACACCAATCCTTGGACCC
HPRT1	NM_000194.1	TGTGATGAAGGAGATGGGAGGCCATCACATTGTAGCCCTCTGTGTCTCAAGGGGGCTATAAATCTTTGCTGACCTGCTGGATTACATCAAAGCACTG
LDHA	NM_005566.1	CAGAATGGAATCTCAGACCTTGTGAAGGTGACTCTGACTTCTGAGGAAGAGGCCCGTTTGAAGAAGAGTGCAGATACACTTTGGGGGATCCAAAAGGAGC
PGK1	NM_000291.2	GCAAGAAGTATGCTGAGGCTGTCACTCGGGCTAAGCAGATTGTGTGAATGGTCCGTGGGGGTATTTGAATGGGAAGCTTTTGCCTGGGAACCAAAGC
POLR1B	NM_019014.3	GGAGAACTCGGCCTTAGAATACTTTGGTGAATGTTAAAGGCTGCTGGCTACAATTTCTATGGCACCAGAGGTTATATAGTGGCATCAGTGGGCTAGAA
POLR2A	NM_000937.2	TTCAAAGAAGCCAAAGACTCCTTCGCTTACTGTCTTCTGTTGGGCCAGTCCGCTCGAGATGCTGAGAGAGCCAAGGATATTCTGTGCCGTCTGGAGCAT
RPL19	NM_000981.3	CCAATGCCGAATGCCAGAGAAGGTCACATGGATGAGGAGAATGAGGATTTTGGCCGGCTGCTCAGAAGATACCCTGAATCTAAGAAGATCGATCGCCA
RDLP0	NM_001002.3	CGAAATGTTTCATTGTGGGAGCAGACAATGTGGCTCCAAGCAGATGCAGCAGATCCGCATGTCCTTTCGCGGGAAGGCTGTGGTGTGATGGGCAAGAA
SDHA	NM_004168.1	TGGAGGGGAGGCTTGGCAGCTGCATTTGGCCTTCTGAGGCAGGGTTAATACAGCATGTGTTACCAAGCTGTTTCCACCAGGTACACACTGTTGCA
TBP	NM_003194.3	CGCCGGCTGTTTAACTTCGCTTCCGCTGGCCCATAGTGATCTTTCAGTGACCCAGCAGCATCACTGTTTCTTGGCGTGTGAAGATAACCCAAAGGAATTG
TUBB	NM_178014.2	TGGTGGATCTAGAACCTGGGACCATGGACTCTGTTGCTCAGTCTTTTGGCCAGATCTTAGACCAGACAACCTTTGATTTGGTCAGTCTGGGCGAGG