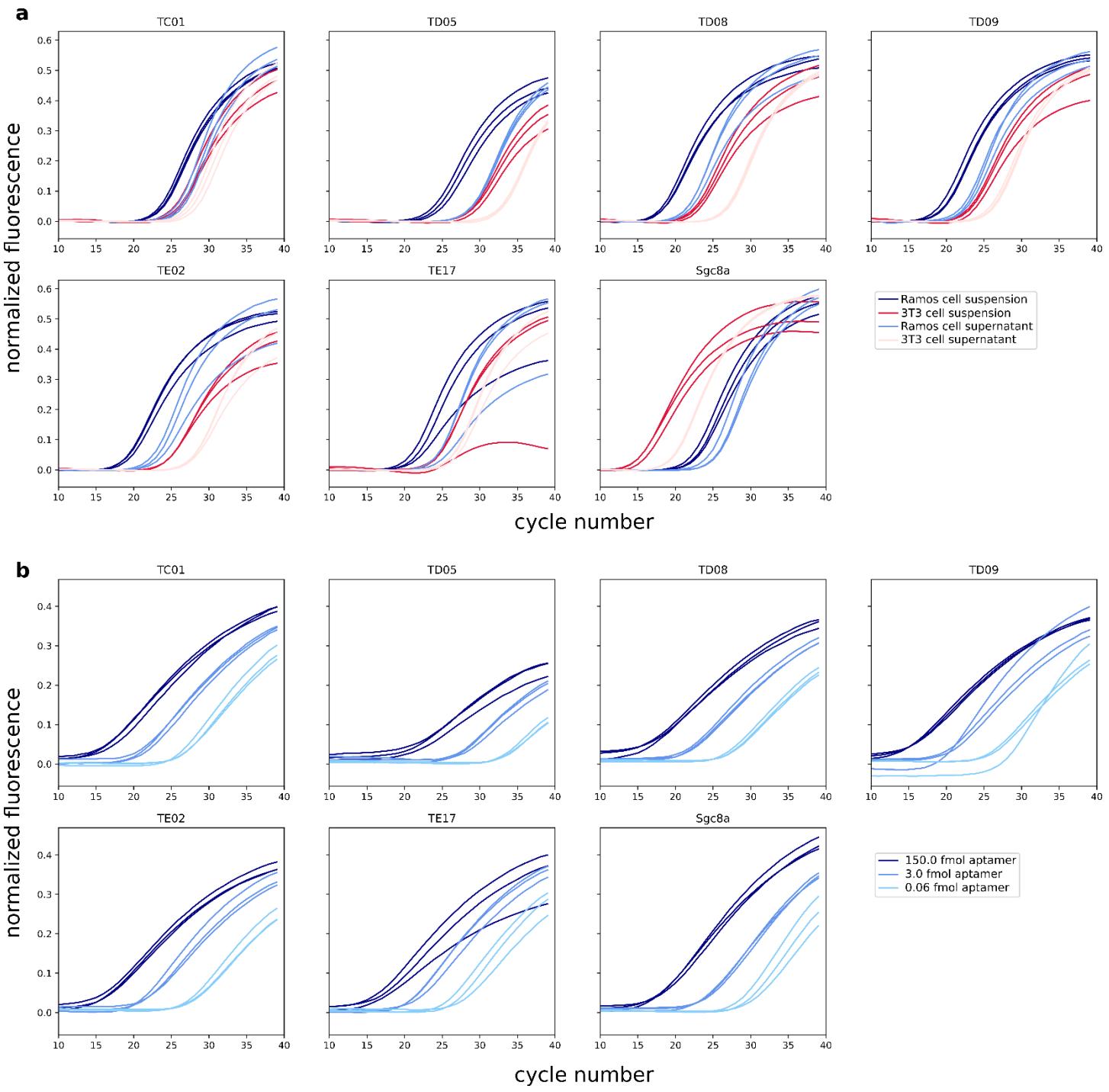


Supplementary Information: Combined aptamer and transcriptome sequencing of single cells

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Description: Supplementary tables and figures



Supplementary Figure 1: qPCR amplification plots of all aptamers modified with a poly-A tail. **a** Association of each aptamer with the two cell lines, Ramos and 3T3 is tested. For each aptamer and both cultures either about 4200 cells in suspension 7 μ l or as background control 7 μ l of supernatant are used as template material. Each curve represents a technical replicate. **b** Corresponding standard curves for each aptamer and associate primer pair is given. In each panel curves represent three independent dilution series.

Supplementary Table 1: Aptamer sequences used in this study. The unique sequence of each aptamer is given in bold face

Name	Sequence
TC01	AAGGAGCAGCGTGGAGGATA ACCAAACACAGATGCAACCTGACTTCTAACGTCAATTGGTG TTAGGGTGTGTCGTCGTGGT BAAAAAAAAAAAAAA
TD05	AAGGAGCAGCGTGGAGGATA AACACC GTTGGAGGATAAGTCGGTGGCTTCAGGGTCTCCTCCGGTG TTAGGGTGTGTCGTCGTGGT BAAAAAAAAAAAAAA
TD08	AAGGAGCAGCGTGGAGGATA CTCTAATTGCCGTATAAGGTCA GGGGTTGGTTGGTCAGGGTCTAGTGC TTAGGGTGTGTCGTCGTGGT BAAAAAAAAAAAAAA
TD09	AAGGAGCAGCGTGGAGGATA GCTGTCATCGGTTATGTGCTTCCAGGGTCCGTGCTGCCGTG TTAGGGTGTGTCGTCGTGGT BAAAAAAAAAAAAAA
TE02	AAGGAGCAGCGTGGAGGATA TAGGCAGTGGTTGACGTCCGCATGTTGGAAATAGCCACGCCT TTAGGGTGTGTCGTCGTGGT BAAAAAAAAAAAAAA
TE17	AAGGAGCAGCGTGGAGGATA CAGCTACGCAATACAAA ACTCCGAACACCTGCTCTGACTGGGTGCTG TTAGGGTGTGTCGTCGTGGT BAAAAAAAAAAAAAA
sgc8a	ATACCAGCTTATTCAATT ACACTTAGAGTTCTAACTGCTGCGCCGCCGGAAA ACTGTACGGTT AGATAGTAAGTGC BAAAAAAAAAAAAAA
sgc3b	ATACCAGCTTATTCAATT CCTGTGGGAAGGCTATAGAGGGGCCAGTCTATGAATA BAAAAAAAAAAAAAA

Supplementary Table 2: Primer sequences

Primer	Sequence
TC/TD/TE nextera adapter	GTCTCGTGGCTCGGAAGGAGCAGCGTGGAGGATA
sgc8a nextera adapter	GTCTCGTGGCTCGGATACCAGCTTATTCAATTACACTTAGAG
sgc3b nextera adapter	GTCTCGTGGCTCGGATACCAGCTTATTCAATTCCGTGGG
TC01 qPCR fw	GCGTGGAGGATAACCAAACAC
TD05 qPCR fw	GCGTGGAGGATAAACACCGTG
TD08 qPCR fw	GCGTGGAGGATACTCTAATTGCC
TD09 qPCR fw	GCGTGGAGGATAGCTGTCATC
TE02 qPCR fw	GCGTGGAGGATAGGCAGTG
TE17 qPCR fw	GCGTGGAGGATACAGCTACG
sgc3b qPCR fw	ATACCAGCTTATTCAATTCCGTGGG
sgc8a qPCR fw	ACCAGCTTATTCAATTACACTTAGAG
TC/TD/TE qPCR rv	ACCACGACGACACACCC
sgc3b qPCR rv	GAGGGGCCAGTCTATGAATA
sgc8a qPCR rv	GCACTTACTATCTAACCGTACAG
DropSeq smart pcr	AAGCAGTGGTATCACGCAGAGT
DropSeq TSO	AAGCAGTGGTATCACGCAGAGTGAATrGrGrG