Supplementary Material

Culture environment regulates amino acid turnover and glucose utilisation in human ES cells

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Table S1. Primer sequences used for qPCR analysis of gene expression

Gene	Forward primer	Reverse primer	
	5'-3'	5'-3'	
B-ACTIN	CGCACCACTGGCATTGTC	TCCTCCTTGATGTCACGCAC	
BRACHYURY	GTGCTGTCCCAGGTGGCTTACAGATG	CCTTAACAGCTCAACTCTAACTACTTG	
GATA4	CTAGACCGTGGGTTTTGCAT	TGGGTTAAGTGCCCCTGTAG	
NANOG	CAAAGGCAAACAACCCACTT	TCTGCTGGAGGCTGAGGT	
OCT4	AGCGAACCAGTATCGAGAAC	TTACAGAACCACACTCGCAC	
SOX2	ATGCACCGCTACGACGTGA	CTTTTGCACCCCTCCCATTT	

Table S2. Measured concentrations of glucose and lactate in control media

	Expected glucose concentration (mM)	Measured glucose concentration (mM)	Expected lactate concentration (mM)	Measured lactate concentration (mM)
mTeSR	13.7	15.2 ± 0.4	0	0
mTeSR + FCS		13.4 ± 0.5		2.4 ± 0.1
mTeSR + BMP4	13.7	14.5 ± 0.3	0	0
mTeSR + KOSR		13.7 ± 0.4		0

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Table S3. Measured concentrations of amino acids in mTeSR and mTeSR supplemented with 20% Life Technologies FCS, compared to the expected concentrations from the published formulation of the medium ¹⁸

** p<0.01; significance was tested for those amino acids showing an increased concentration in serum

containing medium

	mTeSR formulation (μM)	mTeSR (μM)	mTeSR + FCS (μM)
Ala	137	158 ± 3.54	326 ± 4.6**
Arg	548	459 ± 8.5	328 ± 5.6
Asn	137	129 ± 2.1	103 ± 1.5
Asp	137	157 ± 3.4	144 ± 2.8
Glu	137	157 ± 2.9	$301 \pm 3.6**$
Gln	2940	2362 ± 38.1	1973 ± 38
Gly	294	269 ± 7.9	$336 \pm 10.7**$
His	118	151 ± 1.9	145 ± 2.5
Ile	326	318 ± 6.1	291 ± 3.5
Leu	354	352 ± 7.1	333 ± 3.8
Lys	391	358 ± 5.5	331 ± 3.2
Met	90.6	89 ± 1.9	76 ± 1.2
Phe	169	187 ± 3.5	182 ± 2.1
Pro	216	233 ± 2.8	237 ± 2.1
Ser	294	304 ± 8.9	300 ± 8.1
Thr	352	343 ± 6.4	301 ± 3.2
Trp	34.6	68 ± 1.2	71 ± 1.3
Tyr	168	174 ± 3.1	160 ± 1.7
Val	355	355 ± 6.4	355 ± 3.8

Table S4. Measured concentrations of amino acids in mTeSR and mTeSR supplemented with 20% Life Technologies KOSR, compared to the expected concentrations from the published formulation of mTeSR 18 and mTeSR + KOSR

Expected concentrations for amino acids in KOSR were taken from the preferred embodiment of the supplement 20

	mTeSR formulation (μM)	mTeSR (μM)	mTeSR + KOSR formulation (μM)	mTeSR + KOSR (μM)
Ala	137	144 ± 2.5	109.6	123 ± 4.9
Arg	548	581 ± 4.9	438.4	463 ± 16.1
Asn	137	142 ± 2.2	109.6	113 ± 2.7
Asp	137	139 ± 8.8	109.6	106 ± 4
Glu	137	138 ± 6.2	109.6	105 ± 6
Gln	2940	2614 ± 29.5	2352	2065 ± 46.5
Gly	294	296 ± 4.5	1176.4	621 ± 19.1
His	118	116 ± 7.6	1667	925 ± 36.3
Ile	326	332 ± 6.4	6512	4957 ± 118.8
Leu	354	358 ± 6.8	283.2	369 ± 9.4
Lys	391	393 ± 10.7	312.8	308 ± 8.3
Met	90.6	81.6 ± 4.3	465.68	172 ± 4.4
Phe	169	188 ± 4.1	2847.2	2129 ± 55.6
Pro	216	242 ± 5.9	7111.2	6445 ± 144.5
Ser	294	281 ± 4.7	2290.6	1635 ± 46.5
Thr	352	353 ± 9	5038.2	3452 ± 82
Trp	34.6	47 ± 2.9	812	455 ± 13
Tyr	168	226 ± 4.9	752	225 ± 4.9
Val	355	370 ± 7.4	5450	3977 ± 92.6

Table S5. Proliferation of human ES cells, shown as the % of cells positive for phosphorylated histone H3, in medium conditions used, and final live cell density, estimated be Alamar Blue staining, shown as fluorescent intensity

P-value determined by Student's t-test when compared to cells cultured in mTeSR. Analysis was of 20 fields from two independent replicates (proliferation rate) and 6 replicate wells (final live cell density)

	% cells in mitosis ± s.e.m.	P-value	Fluorescent intensity (relative units) ± s.e.m.	P-value
mTeSR	3.6 ± 0.3		2.95 ± 0.3	
mTeSR + F12	3.6 ± 0.3	0.86	2.76 ± 0.2	0.65

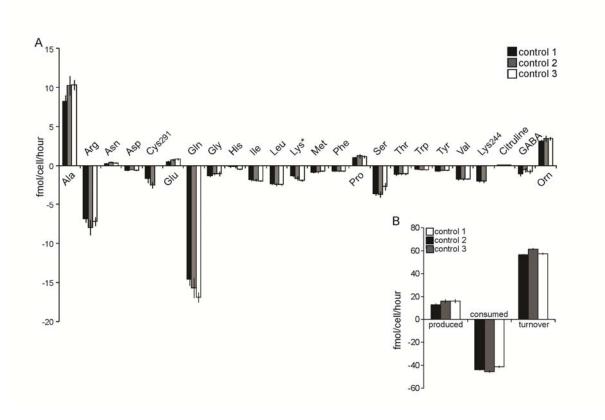


Fig. S1. Comparison of amino acid use in human ES cell cultures across experimental procedures. (A) Spent medium from the final 24 h of culture of human ES cells in mTeSR1 for 7 days was analysed for the presence and concentration of amino acids. Amino acid production or consumption was normalised to cell number and is expressed as fmol/cell/h. n = 8 (Control 1); n = 6 (Control 2); n = 12 (Control 3). Error bars represent s.e.m. Data were analysed using ANOVA, *P < 0.05. (B) Total amino acid production, consumption and turnover by cells cultured in mTeSR1.

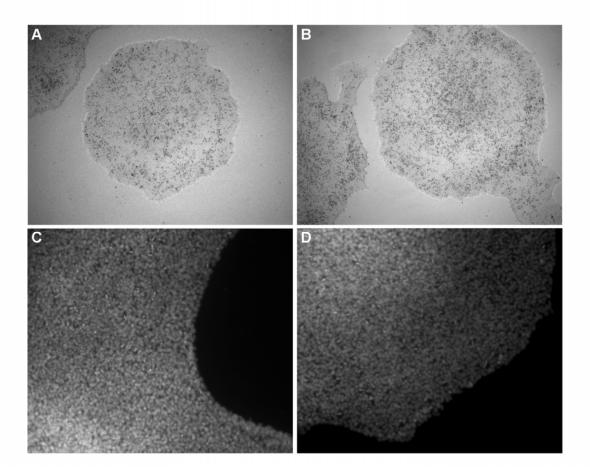


Fig. S2. Morphology of human ES cells cultured in mTeSR1 supplemented with 20% Hams F12 medium. Human ES cells were cultured for 3 days in mTeSR1 before being changed into mTeSR1 (A, C) or mTeSR1 supplemented with 20% Hams F12 (B, D) and maintained for a further 4 days. Images of colon morphology were captured in phase contrast at $4 \times$ magnification (A, B) or with fluorescence microscopy after staining for DNA with DAPI (C, D; $20 \times$ magnification). n = 3; representative images are shown.