

SUPPLEMENTARY TABLE S2. CELL CHARACTERIZATION

<i>Passage number</i>	<i>Day</i>	<i>Group</i>	<i>CD34</i> (-), %	<i>CD45</i> (-), %	<i>CD73</i> (+), %	<i>CD90</i> (+), %	<i>CD105</i> (+), %	<i>STRO-1</i> (+), %
P2-5	10	Untreated	99.58±0.34	86±14	4.0±2.7	92.0±5.6	1.2±1.4	0.31±0.29
		<i>MATH-1</i>	96.2±7.6	84±14	10±10	77±40	2.3±3.1	4.1±6.4
		<i>HATH1</i>	98.8±2.0	83±19	4.9±3.4	93.1±8.5	0.50±0.70	0.46±0.90
		<i>H1/H5</i>	99.74±0.49	89±14	2.3±1.7	93.9±6.4	0.06±0.07	0.03±0.02
		<i>MATH-1/H1/H5</i>	99.4±1.2	80±23	4.0±4.4	81±34	0.10±0.14	0.13±0.17
		<i>HATH1/H1/H5</i>	97.8±4.8	86±10	8.5±15	93.6±7.1	0.10±0.07	0.09±0.16

Values are presented as means with standard deviations. P2-5 indicates that cells were thawed at P2 and subcultured to P5 for use in experiments. The values are representative of cells from five different umbilical cords ($n=5$).