

Gene Name	Description	Studies	Studies with FC	Total Sample Sizes	Total Sample Sizes with FC	Mean Fold Change	Range
KRT8	Keratin 8	3 [19,45,60]	3	38	38	-3.45	-5.14 to -2.36
2 Studies - Greatest sample size							
SEPP1	Selenoprotein P, plasma, 1	2 [6,19]	2	37	37	-10.67	-18.00 to -3.33
IL12RB1	Interleukin 12 receptor, beta 1	2 [8,45]	2	40	40	-2.70	-3.33 to -2.06
CETP	Cholesteryl ester transfer protein, plasma	2 [6,8]	2	51	51	-2.67	-3.33 to -2.00
ITPKB	Inositol 1,4,5-trisphosphate 3-kinase B	2 [6,8]	2	51	51	-2.50	-3.33 to -1.67
2 Studies - Lowest/Moderate sample size							
CA2	Carbonic anhydrase II	2 [8,19]	2	22	22	-16.50	-28.00 to -5.00
CEACAM1	Carcinoembryonic antigen-related cell adhesion molecule 1 (biliary glycoprotein)	2 [8,19]	2	22	22	-15.00	-27.5 to -2.50
GPA33	Glycoprotein A33 (transmembrane)	2 [8,19]	2	22	22	-12.50	-21.67 to -3.33
SLC26A3	Solute carrier family 26, member 3	2 [8,19]	2	22	22	-12.17	-21.00 to -3.33
FKBP1A	FK506 binding protein 1A, 12kDa	2 [8,19]	2	22	22	-10.00	-15.00 to -5.00
KRT17	Keratin 17	2 [8,19]	2	22	22	-6.58	-9.83 to -3.33
THBS2	Thrombospondin 2	2 [8,60]	2	30	30	-5.70	-8.90 to -2.50
PRSS8	Protease, serine, 8 (prostasin)	2 [8,19]	2	22	22	-5.65	-8.80 to -2.50
KIAA0828	Adenosylhomocysteinase 3	2 [8,19]	2	22	22	-4.54	-5.75 to -3.33
ACADS	Acyl-Coenzyme A dehydrogenase, C-2 to C-3 short chain	2 [8,19]	2	22	22	-4.33	-5.33 to -3.33
UQCR	Ubiquinol-cytochrome c reductase, 6.4kDa subunit	2 [8,19]	2	22	22	-3.36	-4.22 to -2.50