Supplementary figures

Targeted and untargeted proteomics of plasma from cord blood with high vs low hematopoietic stem and progenitor cell count

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{r, setup, include=FALSE} # knitr::opts_knit$set(root.dir =
'/Users/xrydbh/OneDrive - University of Gothenburg/Neored') #
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Box plots MS

Fig.S1

Fig.S2

Significant_pathways_in_pw_hierarchy

Venn diagram of pathway overlap

Mass spectrometry 32 with narrow distribution out of 44 proteins with p < 0.05; FC > 2 AHNAK_Q09666 AHSP_Q9NZD4 ALOX12B_O75342 ANXA1_P04083 ANXA2_P07355 APOC2_P02655 ARG1_P05089 CA1_P00915 CAST_P20810 COL1A1_P02452 CTNNB1_P35222 Median centred log2 normalized abundance IMPA2_O14732 FLG2_Q5D862 GSDMA_Q96QA5 H1F0_P07305 KPRP_Q5T749 KRT1_P04264 3 2 1 0 KRT6A_P02538 LEAP2_Q969E1 LOC101060545_P49247 MUC7_Q8TAX7 PLEC_Q15149 S100A14_Q9HCY8 3 1 0 -1 SCARB2_Q14108 SEC61A1_P61619 SERPINB12_Q96P63 SERPINB4_P48594 SH3BGRL3_Q9H299 UBA52_P62987 3 1 0 -1 -2 High Low High Low High Low . High Low VAV2_P52735 VWF_P04275 3 2 1 0

Figure 1: Fig.S1 Box plots of proteins in the MS data with >two-fold abundance differences and p<0.05 between groups (CD34 high vs CD34 low).

Top table protein

. High

Low

High

Low

Mass spectrometry 12 with wider distribution out of 44 proteins with p < 0.05; FC > 2

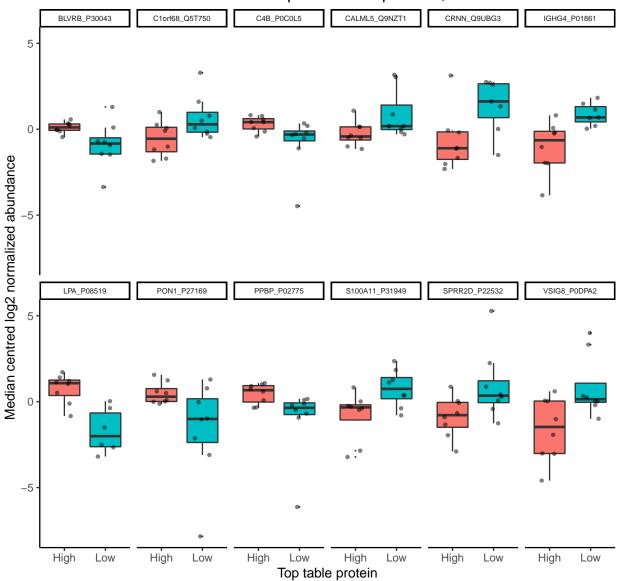


Figure 2: Fig.S1

Olink 7 proteins with p < 0.05; FC > 2 FKBP4_Q02790 HDGF_P51858 HO-1_P09601 LEP_P41159 2 High Low PAI_P05121 PARP-1_P09874 vWF_P04275

Median centred relative NPX (to sample sum)

0

-2

High

Low

Figure 3: Fig.S2

Low

Low High
Top table protein

Low

High

```
Reactome
  i--Transport of small molecules
      °--16 Plasma lipoprotein assembly, remodeling, and clearance (20-98)
          !--1 Plasma lipoprotein remodeling (13-54)
          °--7 HDL remodeling (9-24)
  I--Hemostasis
      "--3 Response to elevated platelet cytosolic Ca2+ (64-146)
          °--2 Platelet degranulation (64-139)
) |
     *--Diseases of signal transduction by growth factor receptors and second messengers
 -
          °--Oncogenic MAPK signaling
: 1
             I--Signaling by high-kinase activity BRAF mutants (13-44)
             :--15 Signaling by RAS mutants (14-54)
             I--12 Signaling downstream of RAS mutants (14-54)
             I--14 Paradoxical activation of RAF signaling by kinase inactive BRAF (14-54)
             °--13 Signaling by moderate kinase activity BRAF mutants (14-54)
: |--Signal transduction
      °--MAPK family signaling cascades
          °--MAPK1-MAPK3 signaling
              °--RAF-MAP kinase cascade
 - 1
                 °--6 MAP2K and MAPK activation (13-49)
---18 Integrin signaling (10-39)
· | |--8 p130Cas linkage to MAPK signaling for integrins (9-22)
      °--17 GRB2:SOS provides linkage to MAPK signaling for Integrins (10-20)
: :--Developmental biology
      °--9 Keratinization (8-226)
         °--10 Formation of the cornified envelope (8-13)
 °--Metabolism of proteins
      °--11 Translation (16-339)
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Figure 4: Fig.S3

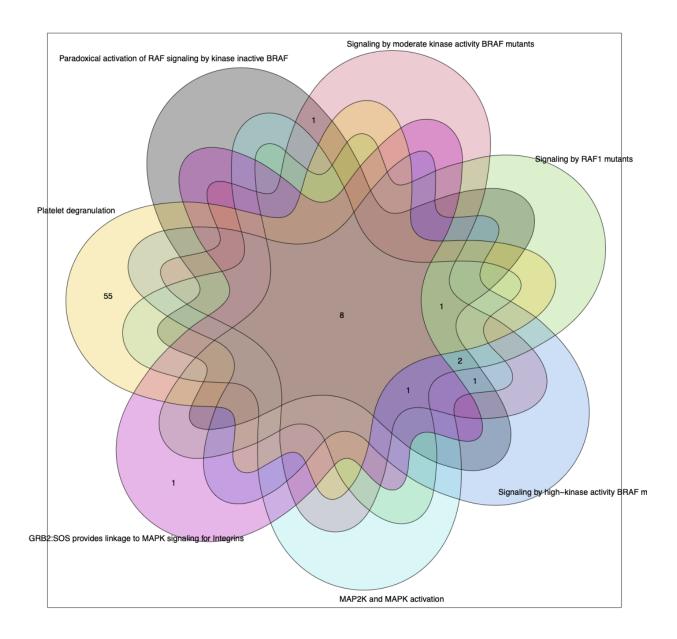


Figure 5: Fig.S4