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') #

## Box plots MS

## Fig.S1

![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).](data:application/pdf;base64,)

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).

![Fig.S1](data:application/pdf;base64,)

Fig.S1

## Fig.S2

![Fig.S2](data:application/pdf;base64,)

Fig.S2

## Significant\_pathways\_in\_pw\_hierarchy

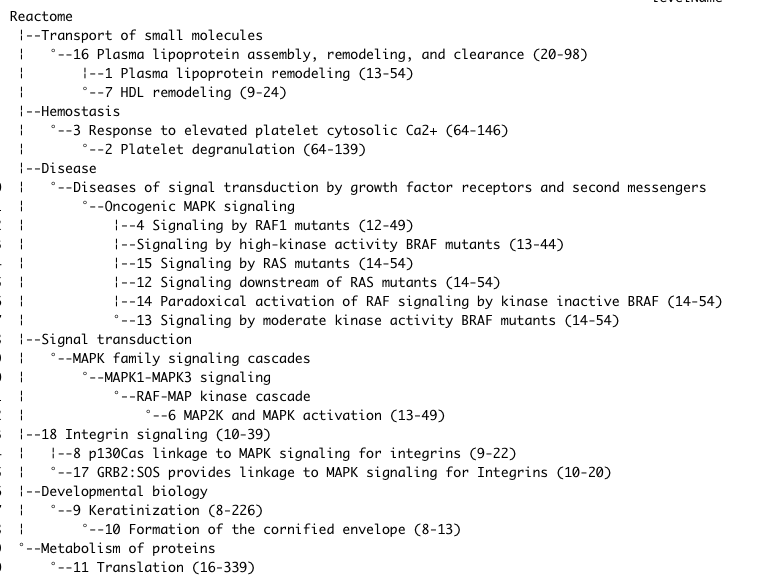


Fig.S3

## Venn diagram of pathway overlap

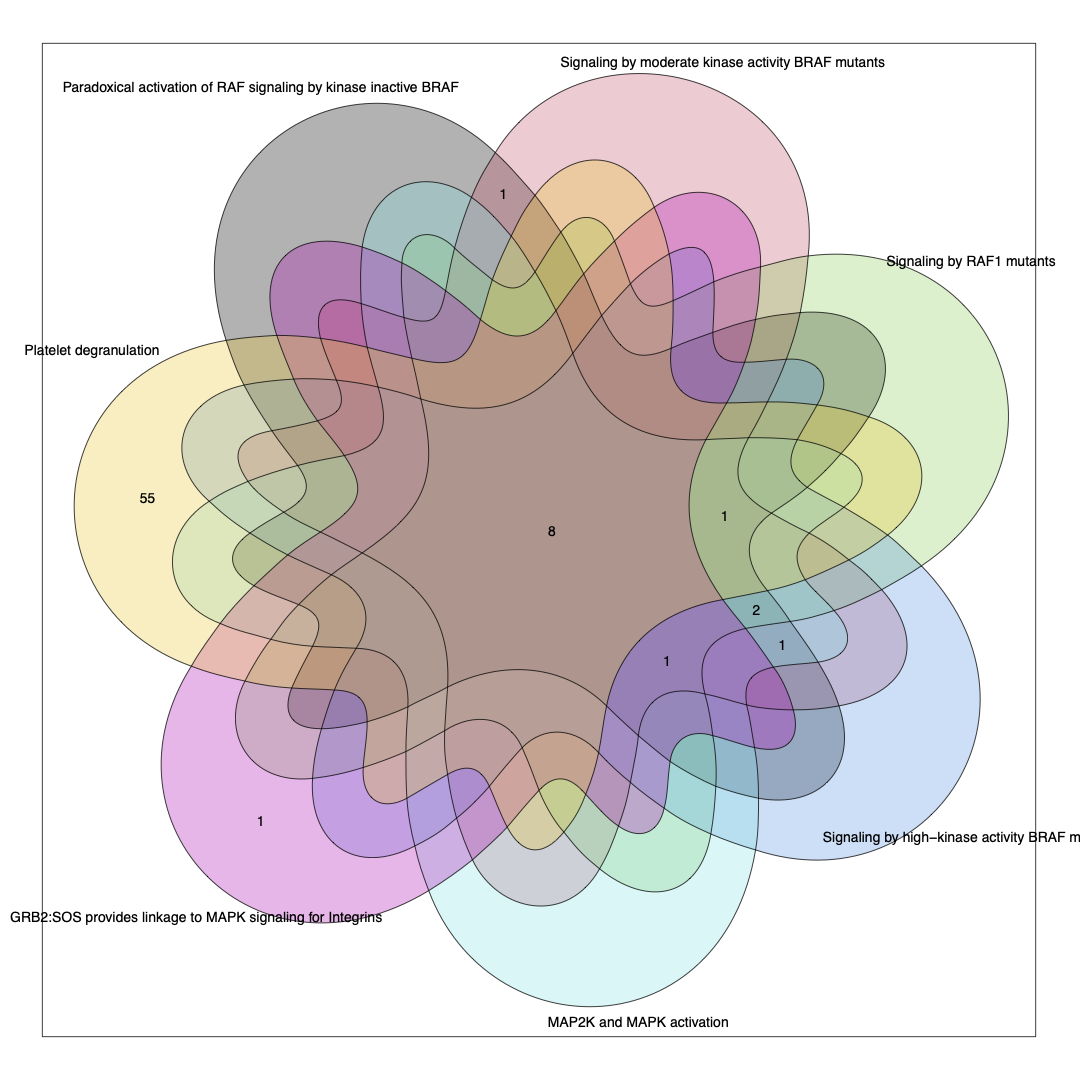


Fig.S4