

**A network analysis of cofactor-protein interactions for analyzing associations
between human nutrition and diseases**

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Supplementary Information (SI)

Table S1. Cofactor-protein interactions and genetic variants in cofactor-binding sites. List of cofactor-protein interactions from all mined databases and genetic variants in cofactor-binding sites from the Uniprot database.

Table S2a. Gene ontology annotation results for each module detected in the cofactor-interacting proteins network.

Table S2b. Biological pathway enrichment results for each module detected in the cofactor-interacting proteins network.

Table S3. Tissue expression of cofactor-interacting proteins.

Table S4. Cofactor-disease interactions. OMIM and GWAS diseases associated with genes encoding for cofactor-interacting proteins.

Figure S1. Cofactor classification.

Figure S2. Overlap of cofactor-interacting proteins between the four different databases.

Figure S3. Per cofactor representation of the degree of connectivity of cofactors-interacting proteins.

Cofactors

Inorganic

Copper (Cu)
Iron (Fe)
Magnesium (Mg)
Manganese (Mn)
Molydenum (Mo)
Nickel (Ni)

Zinc (Zn)
Cobalt (Co)
Potassium (K)

Calcium (Ca)
Chloride (Cl)

Selenium (Se)
Phosphate

Metal complex

Fe-S complex

Organic

Vitamin

Thiamine pyrophosphate [Thiamine (B1)]
Flavin adenine mono/dinucleotide (FMN/FAD)[Riboflavin (B2)]
NAD⁺ and NADP⁺ [Niacin (B3)]
Coenzyme A/Phosphopantetheine [B5]
Pyridoxal phosphate [Pyridoxine (B6)]
Biotin (B7)
Tetrahydrofolic acid [Folic acid (B9)]
Methylcobalamin/AdenosylCobalamine (B12)

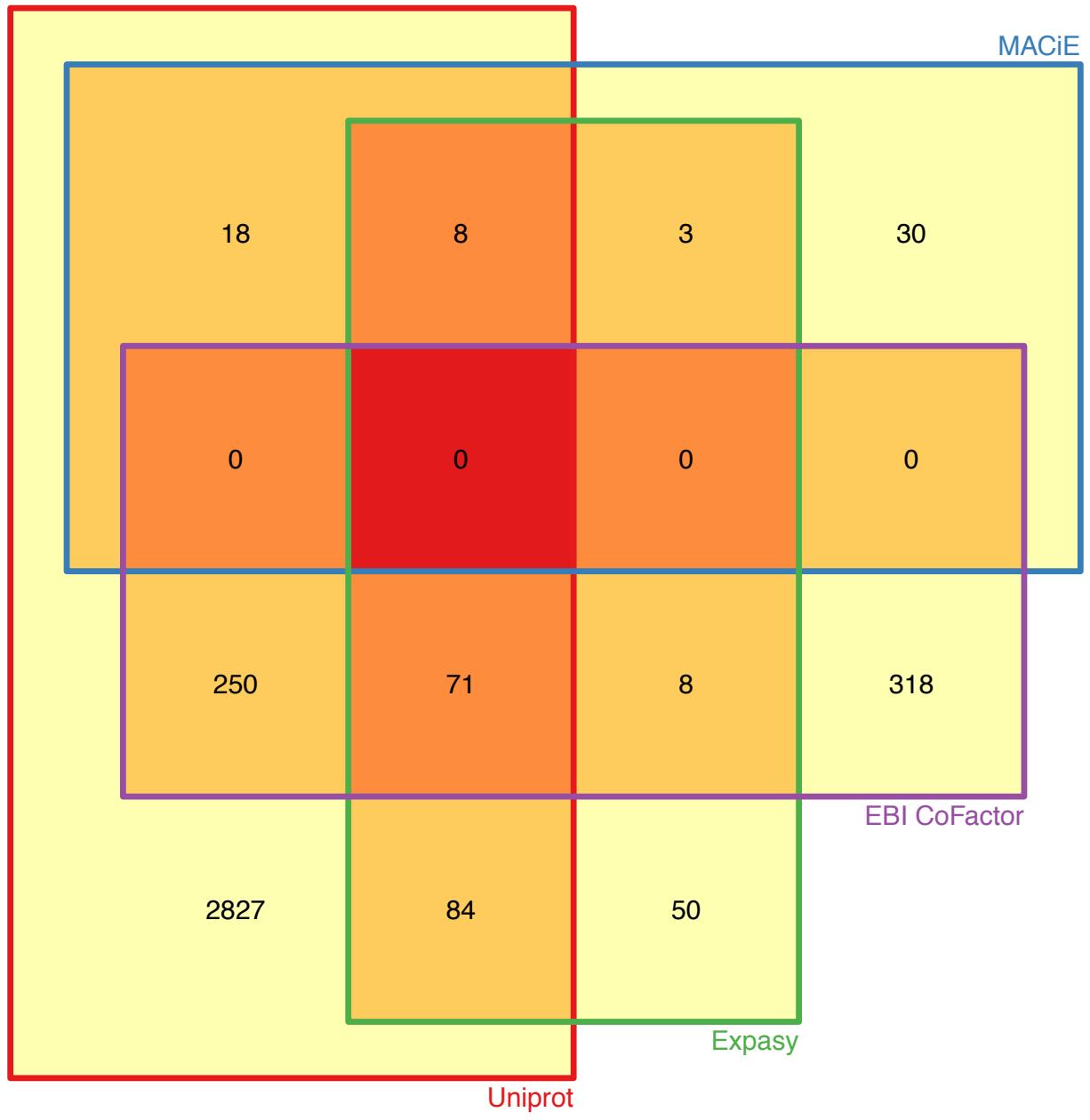
Ascorbic acid (C)
Menaquinone (K)

Transcription factor ligands

Retinol(A): RXR-RAR
D3:VDR
E: PXR

Non-vitamin

In box non-vitamin box
Biopterin
Glutathione (GSH)
Heme
Heme-thiolate
Lipoic Acid (LA)
Molybdopterin (MPT)
Orthoquinone
Pyrroloquinoline quinone (PQQ)
S-Adenosyl methionine (SAM)
S-Adenosyl-L-homocysteine (SAH)
Tetrahydrobiopterin (BH4)
Topaquinone
Ubiquinone (CoQ)



Degree distribution of cofactors

