

Unknown

[illegible]

The diagram illustrates a network of gene and protein interactions, organized into three cellular compartments: Plasma Membrane, Cytoplasm, and Nucleus. Nodes are represented by colored circles (yellow, blue, grey) and rectangles (white, grey). Edges are solid or dashed lines. Some nodes have numerical values next to them.

Plasma Membrane:

- Nodes: **MTN2** (yellow circle, 1.468), **LAMC2** (yellow circle, 1.562), **LAMA3** (yellow circle, 1.431), **ND1** (grey circle, 1.052), **PTHLH** (yellow circle, 1.313), **IL1** (yellow circle, 1.314), **AREG** (yellow circle, 1.495), **MMP1** (yellow circle, 1.730), **ITGA2** (yellow circle, 1.243), **ITGB8** (yellow circle, 1.123), **PCDH7** (yellow circle, 1.508), **EPB41L1** (blue circle, 1.054), **CO24** (yellow circle, 1.069), **IL19** (blue circle, -1.120).

Cytoplasm:

- Nodes: **EGFR** (yellow circle, 1.450), **CO44** (yellow circle, 1.246), **CKCAR** (white rectangle), **G8P1** (yellow circle, 1.081), **PTCS2** (yellow circle, 1.128), **SOCS2** (blue circle, 1.010), **PLCE1** (blue circle, 1.102), **TAGLN** (blue circle, 1.821), **RCST6** (blue circle, -1.302).

Nucleus:

- Nodes: **EGR1** (yellow circle, 3.750), **GADD45A** (yellow circle, 1.692), **TNFAIP3** (grey circle, 1.422), **ELF3** (blue circle, -1.072), **ELF2** (yellow circle, 1.252), **MYOCD** (yellow circle, 1.948), **ID1** (blue circle, -1.083), **ATF3** (yellow circle, 1.383), **RBM39** (blue circle, -1.019).