

# Supplementary Material

## Typical Lung Carcinoids with Metastasis: Potential Role of MicroRNAs in the Regulation of Adaptive Immunity Associated with Disease: a Case Study

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## Supplementary Tables

Due to their size, please use the following links to download and access the Supplementary Tables:

Supplementary Table 1: Complete list of deregulated miRNAs ( $FC \geq 2$  and  $p < 0.05$ ) in tumour and metastasis.

[https://www.cellphysiolbiochem.com/Articles/000320/SM/Supplementary\\_Table\\_1.xlsx](https://www.cellphysiolbiochem.com/Articles/000320/SM/Supplementary_Table_1.xlsx)

Supplementary Table 2: List of miRNA predicted targets (miRTarBase).

[https://www.cellphysiolbiochem.com/Articles/000320/SM/Supplementary\\_Table\\_2.xlsx](https://www.cellphysiolbiochem.com/Articles/000320/SM/Supplementary_Table_2.xlsx)

Supplementary Table 3: Statistically significantly enriched pathways including miRNA target genes.

[https://www.cellphysiolbiochem.com/Articles/000320/SM/Supplementary\\_Table\\_3.xlsx](https://www.cellphysiolbiochem.com/Articles/000320/SM/Supplementary_Table_3.xlsx)

Supplementary Table 4: miRNA target genes have deregulated expression levels in typical vs. atypical carcinoids (GSE35679).

[https://www.cellphysiolbiochem.com/Articles/000320/SM/Supplementary\\_Table\\_4.docx](https://www.cellphysiolbiochem.com/Articles/000320/SM/Supplementary_Table_4.docx)

## Supplementary Figures

**Supplementary Fig. 1.** miRNA-mRNA network enrichment of target genes associated with the adaptive immune response.

