

Supplemental Information

**Table.** Semi-quantitative densitometric analyses indicate that dominant-interfering Pyk2 (AdCRNK) decreases basal and inducible tyrosine phosphorylation of Pyk2 and p130Cas relative to the unstimulated AdGFP cells.

| Tyrosine residue             | Virus<br>(Ad) | Stimulation (minutes) |                  |              |              |              |              |  |
|------------------------------|---------------|-----------------------|------------------|--------------|--------------|--------------|--------------|--|
|                              |               | no stim               | CXCL12 [20ng/ml] |              |              |              | LPA [1μg/ml] |  |
|                              |               |                       | 1'               | 5'           | 10'          | 30'          | 5'           |  |
| pY402 Pyk2                   | GFP           | 1.00                  | 0.84 ± 0.21      | 0.93 ± 0.20  | 1.35 ± 0.18  | 1.06 ± 0.18  | 0.82 ± 0.24  |  |
|                              | CRNK          | 0.58 ± 0.17           | 0.59 ± 0.18      | 0.53 ± 0.04* | 0.49 ± 0.07* | 0.65 ± 0.15  | 0.57 ± 0.03* |  |
| pY410<br>p130 <sup>Cas</sup> | GFP           | 1.00                  | 1.01 ± 0.10      | 1.45 ± 0.39  | 1.33 ± 0.30  | 1.26 ± 0.13  | 0.88 ± 0.03* |  |
|                              | CRNK          | 0.39 ± 0.06*          | 0.74 ± 0.08      | 0.73 ± 0.01* | 0.87 ± 0.45  | 0.70 ± 0.05* | 0.72 ± 0.14  |  |

Data are relative units compared to unstimulated AdGFP control cells. Values are mean ± SEM of 3 representative immunoblot analyses. Asterisk indicates the statistically significant decrease in Pyk2 or p130Cas phosphorylation relative to the unstimulated AdGFP infected cells.