Supplementary Table-1: Different buffers, used during the optimization of phosphopeptides enrichment process. (highlighted in green \rightarrow delivered best results)

| Loading buffers | Washing buffers | Elution buffers |
|-----------------|-----------------------------------|------------------------|
| | | |
| 40%ACN/ 0.1%TFA | 20 mg DHB/ mL / 2%ACN / 0.5 % TFA | 100 mM Imidazole |
| | | |
| 40%ACN/ 1%TFA | 0.2% HCl | 0.5 % NH₄OH |
| | | |
| 40%ACN/ 4%TFA | H ₂ O | |
| | | |
| 40%ACN/ 3%TFA | | |



1400

1200

1600

1800

2000

2200

2400

2600

2800

3000

3200

m/z



Supplementary figure 3:

MALDI mass spectra of phosphopeptide enrichment by Er-IMAC using different loading buffers (A-D). The maximum number of phosphopeptides could be isolated using 40% ACN/3% TFA as a loading buffer (D). DHB matrix. 400 shots. 337 nm laser, Ultraflex I (Bruker).

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Supplementary figure 4: MALDI mass spectra of the different washings buffers. During washing **(A)** and **(B)**, some of the phosphopeptides get lost (marked with red asterisk). A neutral wash with deionized water **(C)** was observed to cause no loss of phosphopeptides. DHB matrix. 400 shots. 337 nm laser, Ultraflex I (Bruker).



Supplementary figure 5: MALDI Mass spectra of two different elution buffers. Only a small number of phosphopeptides can be eluted with 100 mM imidazole (A). Highest number of phosphopeptides could be enriched with 0.5% ammonia solution (B).