Dear Editor,

We would be pleased if you consider our manuscript "Lost in translation: On the

impact of data coding on penalized regression with interactions" for publication in Ge-

netics Selection Evolution.

In recent scientific work on genomic prediction, a tendency towards models incor-

porating gene by gene or gene by environment interactions can be observed [1, 2, 3, 4].

It has been shown that the additive ridge regression best linear unbiased prediction is

not affected by data coding [5, 6], but that models with products of marker values as

additional predictor variables show a sensitivity to changes of the marker coding[7, 6].

In our work, we the investigate the cause of this loss of invariance to changes in

the marker coding, in a general setup of penalized regression on models defined by

polynomials in the predictor variables. Our results are not limited to ridge regression,

but for instance also apply to LASSO or to any other penalized regression, and to any

type of interaction (GxG or GxE) modelled by products of predictor variables or by

covariance matrices given by Hadamard products.

We acknowledge that our considerations are brief, and mainly of theoretical nature.

However we are confident that a journal with focus on quantitative genetics as GSE is

the right place to present our work, whether as a full length paper, or as a short com-

munication, whichever suits best to the Journal.

Best regards,

Johannes Martini

Francisco Rosales

The following persons are suggestions for potential reviewers:

1

- Daniel Gianola, Department of Animal Science, University of Wisconsin-Madison, gianola@ansci.wisc.edu
- Gustavo de los Campos, Department of Epidemiology and Biostatistics, Michigan State University, gdeloscampos@epi.msu.edu
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