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## Correction

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Corrections to ac2030255

*Title of the publication:* Isolation of alpaca anti-hapten heavy chain single domain antibodies for development of sensitive immunoassay

*Authors:* Kim HJ, McCoy MR, Majkova Z, Dechant JE, Gee SJ, Tabares-da Rosa S, González-Sapienza GG, Hammock BD..

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The corrections refer to the Supplementary Information section that can be found at <http://pubs.acs.org/doi/suppl/10.1021/ac2030255>

### Corrections:

Incorrect primer sequences for library construction were included in the supplementary material of the manuscript (Page 3, Lines 10–14).

### *Current version:*

DNA fragments encoding the VHH IgG variable domains were amplified by PCR using the forward primer Alp-Vh-F1 SfiI: CAT GCC ATG ACT GTG GCC CAG GCG GCC CAG KTG CAG CTC GTG GAG TCN GGN GG targeting the framework 1 region; and reverse primers AlpVhh-R1 SfiI: CAT GCC ATG ACT CGC GGC CGG CCT GGC CTC GTG GGG GTC TTC GCT GTG GTG CG and AlpVhh-R2 SfiI: CAT GCC ATG ACT CGC GGC CGG CCT GGC CTC GCC TTG TGG TTT TGG TGT CTT GGG corresponding to the short (IgG3) and long (IgG2) hinge region, respectively.

### *Correct version:*

DNA fragments encoding the VHH IgG variable domains were amplified by PCR using the forward primer Alp-Vh-F1 SfiI: CAT GCC ATG ACT GTG GCC CAG GCG GCC CAG KTG CAG CTC GTG GAG TC targeting the framework 1 region; and reverse primers AlpVhh-R1 SfiI: CAT GCC ATG ACT CGC GGC CGG CCT GGC CAT GGG GGT CTT CGC TGT GGT GCG and AlpVhh-R2 SfiI: CAT GCC ATG ACT CGC GGC CGG CCT GGC CGT CTT GTG GTT TTG GTG TCT TGG G corresponding to the short (IgG3) and long (IgG2) hinge region, respectively.

The original version of forward primer will result in increased rates of random mutations due to the guanosine-rich stretch of nucleotides at 3' end. The original versions of reverse primers will result in a frame shift at the N-terminus of the pIII protein.