PART 6 THE BEAST

Remember that I said verification is relatively easy?

Return of the Kummer: a toolbox for genus 2 cryptography

Maria Corte-Real Santos¹ and Krijn Reijnders²

 University College London maria.santos.20@ucl.ac.uk
 Radboud University, Nijmegen, The Netherlands krijn@cs.ru.nl

Q: Is it faster than 1D or 2D? A: No.

2D 1D-SQlsign

Map the 2^{1000} isogeny from 1D SQIsign over \mathbb{F}_{p^2} to a 2D isogeny over \mathbb{F}_p using Scholten's construction and Costello's isogenies.

Requires tons of work as we now don't do a single "short" 2D-isogeny, but a number of blocks.

So, we developed:

- pairing-based techniques
- efficient basis sampling
- point compression
- curve normalisation



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Q: Is it faster than 1D or 2D? A: No.

Q: Is it much more difficult?

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