INTERMITTENT C1-INHIBITOR DEFICIENCY ASSOCIATED WITH RECESSIVE INHERITANCE: FUNCTIONAL AND STRUCTURAL INSIGHT

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Supplementary figure 1. Complement parameters. Serial plasma levels of C1-INH function (white bars), C1-INH antigen (black bars) and C4 antigen (gray bars) in patient T.M. referred as a percentage of the pooled normal plasma used as reference.


Supplementary figure 2. Analysis of the 96-kd band. SDS- PAGE and immuno-blot of plasma samples revealed with in house chicken polyclonal antibodies designed against the whole molecule (anti total C1-INH, left) or against the 10 C -terminal residues of the intact C1-INH ( anti C-ter C1INH, right), not reacting with the cleaved form. Normal human plasma (NHP), the plasma of two patients heterozygous for the C1-INH Mo mutation (Mo), which have already been reported to contain RCL cleaved C1-INH, and patient T.M. plasma with two different C1-INH concentrations $(\mathrm{R} / \mathrm{C})$ are analysed. Intervening lanes were removed for clarity, but the relative positions of the bands were unaltered. *The intervening lines were cut for clarity purposes.


Supplementary figure 3. Expression of C1-INH proteins in P. pastoris cells. Single colonies (15) were selected and grown in glycerol until $\mathrm{OD}_{600 \mathrm{~nm}}$ was between 2 and 6 and then for 3 days in buffered complex medium with methanol $(0.5 \%, \mathrm{v} / \mathrm{v})$. The culture media and cell pellet were tested for secreted and intracellular C1-INH by means of ELISA. The mean extracellular values were 2.1 $\pm 0.3 \mu \mathrm{~g} / \mathrm{ml}$ and $24.6 \pm 4.4 \mu \mathrm{~g} / \mathrm{ml}$ for secreted $\operatorname{Arg} 378$ Cys and wild-type respectively (upwards triangles); and $0.20 \pm 0.05 \mu \mathrm{~g} / \mathrm{ml}$ and $0.53 \pm 0.18 \mu \mathrm{~g} / \mathrm{ml}$ for intracellular $\operatorname{Arg} 378$ Cys and wild-type, respectively (downwards triangles). (WT and R378C for wild-type, open symbols, and Arg378Cys, closed symbols, respectively). P values from unpaired Student's t test are indicated.

| Protease | [Protease $](\mathrm{nM})$ | Substrate | $[$ Substrate $](\mu \mathrm{M})$ | $[$ C1-Inh $](\mathrm{nM})$ |
| :--- | :--- | :--- | :--- | :--- |
| C1s | 1 | H-D-KGR-pNA | 240 | $10-15-20-25-30-35-40$ |
| Kallikrein | 0.025 | H-D-PFR-pNA | 250 | $10-20-30-50-80-130-160$ |

Supplementary Table I. Reagent concentrations for kinetic assays.

| Protease | Wild-type rhC1-Inh | R378C rhC1-Inh |
| :--- | :--- | :--- |
|  | $\mathrm{k}_{\text {inh }}\left(\mathrm{M}^{-1} \mathrm{~s}^{-1}\right)$ | $\mathrm{k}_{\text {inh }}\left(\mathrm{M}^{-1} \mathrm{~s}^{-1}\right)$ |
| C1s | $(4.03 \pm 0.15) \times 10^{4}$ | $(2.95 \pm 0.09) \times 10^{4}$ |
| Kallikrein | $(11.52 \pm 0.50) \times 10^{3}$ | $(6.67 \pm 0.39) \times 10^{3}$ |

Supplementary Table II. Apparent second-order rate constants of inhibition for the interaction between recombinant C1-INH and proteases.

