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Center for Macromolecular Crystallography University of Alabama in Birmingham

Dr. Manuel Navia Senior Scientist Vertex Pharmaceuticals,, Inc.

Refinement of Porcine Pancreatic Elastase Using Data from Crystals Grown in Microgravity

Porcine pancreatic elastase (PPE) crystals grown under microgravity conditions on mission STS-26 of the space shuttle "Discovery" have been shown to diffract to considerably higher resolution than the best PPE crystals grown by us on the ground (DeLucas *et. al.* (1989) *Science* **246:** 651-654). We have now independently refined both the microgravity and ground-based data. Preliminary results of these refinements are summarized below:

| | EARTH GRAVITY | MICRO-GRAVITY. | | |
|----------------|----------------------------|------------------|--|--|
| D-MIN | # of SHELL R- | # of SHELL R- | | |
| RES. to | RFLS FACTOR | RFLS FACTOR | | |
| 3.00A | 3546 19.0% | 3624 14.2% | | |
| 2. 5 0Å | 3044 18.8% | 322 15.0% | | |
| 2.00Å | 5937 18.7 <i>%</i> | 6971 14.8% | | |
| 1.90Å | 1751 20.0% | | | |
| 1.80Å | 1873 22.1% | 4920 15.7% | | |
| 1.70Å | 1990 26.4% | – | | |
| 1.65Å | 944 32.0% | | | |
| 1.60Å | | 6636 17.2% | | |
| 1.40Å | | 8387 20.2% | | |
| 1.30Å | | 2079 26.6% | | |
| | EARTH GRAVITY | MICRO-GRAVITY. | | |
| | # of SHELL R- | # of SHELL R- | | |
| | RFLS FACTOR | RFLS FACTOR | | |
| TOTAL | 19085 19.9% | 35841 15.9% | | |
| Note: Deviatio | ons from ideal bond length | s for both | | |

refinements was appx. 0.020Å.

These results show nearly a doubling of experimental diffraction data for this structure, exceeding 1.3Å resolution. Improved phase information derived from the refined structure of PPE based on this microgravity data has allowed us to interpret previously-uninterpretable electron density obtained from ground-based crystals of a complex of PPE with a chemically-reactive inhibitor. Intermediate stages in the enzyme-inhibitor reaction mechanism in the crystal can now be directly observed. Further refinement of PPE structures is in progress.

| PROTEIN | a (Å) | b (Å) | c (Å) | angles (deg.) | Soln. |
|---|----------------------------------|----------------------------------|----------------------------------|---|--|
| Anhydro-PPE A-PPE + peptide PPE (gravity) PPE (micro-grav) | 50.74 50.94 51.00 50.88 | 57.94 57.91 58.08 58.02 | 75.28 75.33 75.29 75.35 | 90. 90. 90. 90. 90. 90. 90. 90. 90. 90. 90. 90. 90. 90. 90. | SUL SUL SUL SUL |
| Table 3: | | | | | |
| PROTEIN | # reflections | resolution | R-Factor | Bond length deviation (Å) | detector type |
| Anhydro-PPE A-PPE + pepide PPE (gravity) PPE (micro-grav) | 17564 16035 19085 35841 | 1.65Å 1.80Å 1.65Å 1.30Å | 15.7% 16.8% 19.9% 15.9% | 0.019Å 0.016Å 0.016Å 0.016Å | diffractometer diffractometer area detector area detector |

Table 2: