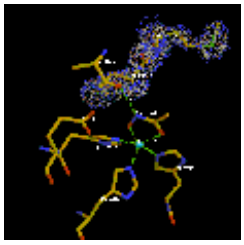
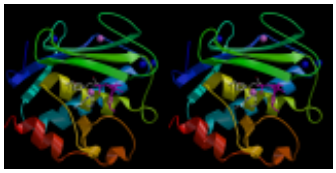
MATRIX METALLOPROTEINASES

Bertini, I., Calderone, V., Fragai, M., Luchinat, C., Mangani, S., Terni, B. **X-Ray structures of binary and ternary enzyme-product-inhibitor complexes of matrix metalloproteinases** Angew.Chem.Int.Ed., 42: 2673-2676, 2003	 PDB code: 1OS2, 1OS9
Bertini, I., Calderone, V., Fragai, M., Luchinat, C., Mangani, S., Terni, B. **Crystal structure of the catalytic domain of human matrix metalloproteinase 10** J.Mol.Biol., 336: 707-716, 2004	 PDB code: 1Q3A
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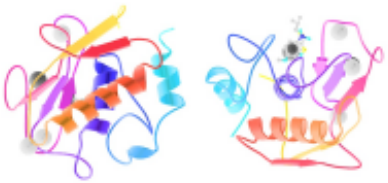
Bertini, I., Calderone, V., Cosenza, M., Fragai, M., Lee, Y.-M., Luchinat, C., Mangani, S., Terni, B., Turano, P. **Conformational variability of matrix metalloproteinases: beyond a single 3D structure** Proc.Natl.Acad.Sci.USA, 102: 5334-5339, 2005	 PDB code: 1Y93, 1RMZ
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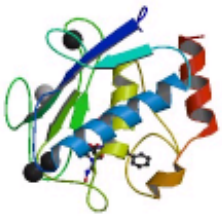
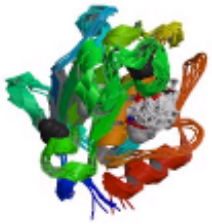
Mannino, C., Nievo, M., Machetti, F., Papakyriakou, A., Calderone, V., Fragai, M., Guarna, A. **Synthesis of bicyclic molecular scaffolds (BTAa): an investigation towards new selective MMP-12 inhibitors** Bioorg.Med.Chem. 14: 7392-7403, 2006	 PDB code: 2HU6
Alcaraz, L.A., Banci, L., Bertini, I., Cantini, F., Donaire, A., Gonnelli, L. **Matrix metalloproteinase-inhibitor interaction: the solution structure of the catalytic domain of human matrix metalloproteinase-3 with different inhibitors** J.Biol.Inorg.Chem. 12: 1197-1206, 2007	 PDB code: 2JNP, 2JT5, 2JT6
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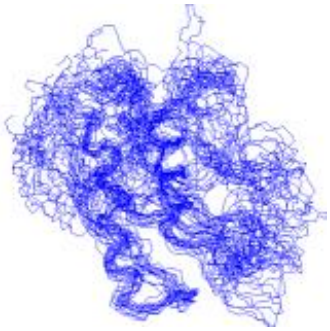
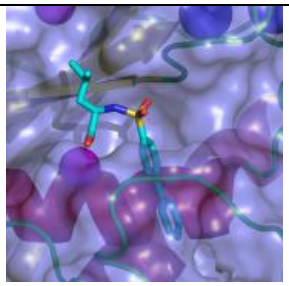
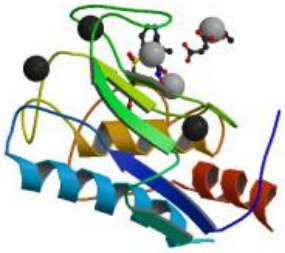
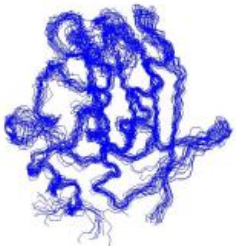
Balayssac, S., Bertini, I., Bhaumik, A., Lelli, Luchinat, C., **Paramagnetic shifts in solid-state NMR of proteins to elicit structural information** Proc.Natl.Acad.Sci.USA, 105: 17284-17289, 2008	 PDB code: 2K9C
Dragoni, E., Calderone, V., Fragai, M., Jaiswal, R., Luchinat, C., and Nativi, C. **Biotin-tagged probes for MMP expression and activation: design, synthesis and binding properties** Bioconjug.Chem., 4: 719-727, 2009	 PDB code: 3EHX, 3EHY
Isaksson, J., Nyström, S., Derbyshire, D., Wallberg, H., Agback, T., Kovacs, H., Bertini, I., Giachetti, A., Luchinat, C. **Does a fast nuclear magnetic resonance spectroscopy- and X-ray crystallography hybrid approach provide reliable structural information of ligand-protein complexes? A case study of metalloproteinases** J.Med.Chem., 52: 1712-1722, 2009	 PDB code: 2W0D
Bertini, I., Bhaumik, A., De Paepe, G., Griffin, R. G., Lelli, M., Lewandowski, J., and Luchinat, C. **High-resolution solid state NMR structure of a 17.6 kDa protein** J.Am.Chem.Soc., 132: 1032-1040, 2010	 PDB code: 2KRJ
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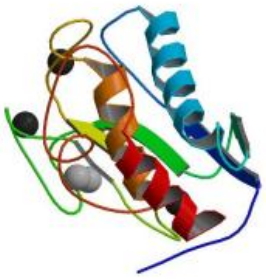
Bertini, I., Calderone, V., Cerofolini, L., Fragai, M., Geraldes, C. F. G. C., Hermann, P., Luchinat, C., Parigi, G., and Teixeira, J. M. C. **The catalytic domain of MMP-1 studied through tagged lanthanides** FEBS Lett., 586: 557-567, 2012	 PDB code: 3SHI
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