Formulae were given for the X-ray diffraction patterns of helical structures, and evidence was provided that the structure of a synthetic polypeptide was based on the α-helix of Pauling and Corey. [The SCI® indicates that this paper has been cited in over 375 publications since 1955.]

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This work was done 35 years ago! In 1951 I had just received my first tenured appointment as a physics lecturer in Cambridge. Francis Crick was a research student in the Medical Research Council Unit of the Cavendish—he had come late into physics research. Vladimir Vand was a research fellow in Glasgow, whom I had met but did not know well. For me the story began when, without much expectation of being able to interpret them, I took some X-ray photographs of a specimen of poly-y-methyl-L-glutamate that my professor, Sir Lawrence Bragg, had obtained from another research group. My diary for October 2, 1951, notes: “The Prof shows a touching faith in my ability to extract a complicated crystal (sic) structure from almost no data”—the material was only semicrystalline, and I was used to looking at crystals. A month or so later Bragg received, possibly as a referee, a paper by Vand on the theory of X-ray diffraction by helical structures. He passed it on to me, and I concluded that Vand’s answer was correct for a continuous helix but not for atoms on a helix. Crick also saw the paper, and when we compared notes the following day, we found that we had arrived at the same (correct) answer by different routes. A few days later I suddenly realised that the photographs of poly-y-methyl-L-glutamate, which I had put aside, could be explained as the diffraction pattern of atoms on helices of different radii. The structure turned out to be based on the α-helix of Linus Pauling and Robert B. Corey. It was, I believe, the first fairly conclusive experimental evidence for the existence of a helical structure at the molecular level, and Crick and I published a short note on the subject.

The main value of this work, seen in retrospect, is that it was a first step on the road to the discovery of the structure of DNA by Jim Watson and Crick. The first I knew of that work was when Crick arrived excitedly in my office to take me to see the model of a double helix that they had built. Actually, it was their first and incorrect version, but I would not have been more impressed by the correct version—I had not seen the experimental evidence contained in photographs of B-type DNA, and I distrusted “speculation.” It was some time before I gradually became convinced of their tremendous success. [The α-helix has now been recognized as a feature of the majority of protein structures, and a vast industry of protein crystallography has developed.]

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