Iron Alum as a Mordant in Cytology of Uredinales

The cytology of fungi has remained relatively unexplored, as the vegetative nuclei are too small to define with precision details of nuclear structure and divisional figures. Information on these is primarily based on the study of meiotic division of the diploid nucleus during germination of the resting spore (zygote cell). The chromosomes often show a weakly staining reaction with the usual aceto-carmine methods and seldom stain deep enough for analytical study of configurations of the chromatin material. Addition of a ferric salt hardly improves the stain density. Thus, choice of suitable stains and staining techniques to bring out critical details has been an important factor, as the stain reactions have often been incomplete, uncertain or erratic. Use of iron alum as a mordant is well known in the advanced and elaborate procedures for staining the nuclei in the higher plant cells. However, not infrequently these have been unsatisfactory and inadequate for critical cytological studies in fungi and other microorganisms. Interpolation of iron alum solution of a suitable strength for staining the nuclei in Chlorophyceae-Conjugales and Volvocales and in Rhodophyceae3 has been successfully employed in the acetocarmine squash technique. A similar procedure briefly described here proved very satisfactory in the cytological studies of the rust fungi Uredinales as well.

The teliospores were stuck onto the slides by alternately wetting and drying the smears and the slides inverted over wet cotton towelling for moisture condensation as described by Thirumalachar et al. 4,5. The spore germination time varied with different rust species depending on the dormancy period. Germinating teliospores at a desired stage were fixed in situ in modified Carnoy's fixing fluid (6 rectified alcohol: 4 glacial acetic acid: 4 chloroform) for 1 h and transferred directly to 70% alcohol. The slides bearing the spores were brought down to distilled water through 4 descending alcohol concentrations with 5 min in each grade and washed thoroughly with 2–3

changes in distilled water. Immersion of the smears in 0.5% aqueous Taka-diastase (Parke Davis (India) Ltd., Bombay) for 10 min helped clarify the cytoplasm⁶. A dip in distilled water removed the enzyme residue over the smears and they were upgraded back to 70% alcohol through the same alcohol series. The slides may be retained at this stage for any length of time desired without impairing the stainability of the nuclei. The slides were directly transferred to the mordant 0.5% aqueous iron alum for 2-5 min. Concentration of the mordant and time for mordanting would vary with the fungal species. They were thoroughly washed in distilled water. Care was taken not to leave any traces of the mordant on the smears which crystallized the stain or stained the nuclei dark, masking structural details. The spore mounts were flooded with 0.5% carmine in 45% glacial acetic acid and covered as usual. Immediate observation facilitated analysis of nuclear details as the stain darkened gradually with lapse of time. The technique yielded excellent results with a variety of other fungi as well.

Zusammenfassung. Es wird ein Beizverfahren beschrieben, das sich bei der Färbung von Kernteilungsfiguren von keimenden Rotpilzsporen bewährt hat.

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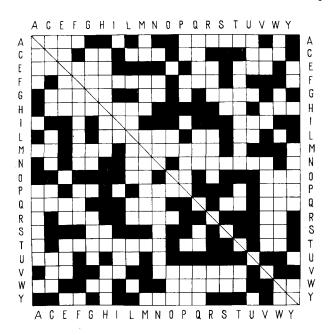
Table of Replaceable Amino Acids in Proteins

In looking for general laws governing the development of primary protein structure¹, some amino acids in peptide chains were found to be frequently replaceable². The regularities observed were expressed in the form of a chart of amino acid interchanges³ (summarized in ⁴ and ⁵). On this basis it was possible, as early as 1962, to construct 8 variants of codons⁶ of the first two bases in triplets, including their sequence. Variant No. 6 (8) (Table I, ref. ⁶) corresponds to the presently accepted codons, with the exception of that for methionine.

At present it is possible to make exact the previously published tables of amino acid interchanges 3,5,6 and to extend them on the basis of existing knowledge of the nucleotide sequence in a degenerated code 7,8. A practical form in this connection is that of a chessboard, as shown in the Table. It is constructed on the assumption that in codon triplets with a sequence 7,8 only a single base will be interchanged for another. The Table differs from the spatial design of Versteeg and Vliegenthart 9

not only in the discrepancies in a number of interchanges, since more recent codons are being used here, but also in the ease of application. The chessboard form makes it possible to find readily all the interchanges of a given

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Full squares indicate amino acids which can be interchanged in the structure of proteins, provided only one single base is changed for another base in the triplet codon.

A list of symbols1 for amino acids

A – asp	G – glu	M - met	Q - gln	U – Ieu
C - cys	H - his	N - asn	R - arg	V – val
$\mathbf{E}-thr$	I - lys	O – tyr	S - ser	W – ile
F - phe	L – ala	P – pro	T - trp	Y - gly

amino acid on a single line, whereas in a three-dimensional scheme⁹ an amino acid must be looked for at different places (e.g. 9 Asn, Leu, Lys, Ser). In consequence of the existence of ApGpU and ApGpC codons⁸ for lysine the Table includes also the interchange Arg/Lys which was not permitted by previously accepted codons. If the codon ApGpA introduced by Nishimura et al. 10 for arginine were included in the Table this would be a further confirmation of the Arg/Lys interchange, but the Table would then contain new interchange possibilities such as Met/Arg, Thr/Arg, and at the same time the interchanges Arg/Ser, Gly/Ser, Lys/Ser and Met/Ser would disappear.

Zusammenfassung. Auf Grund vorausgegangener Experimente, besonders über degenerierte Triplettcodons, wird ein Verfahren entwickelt, aus dem sich die Austauschmöglichkeiten von Aminosäuren in Eiweißstoffe ergeben, dann nämlich, wenn im Codon nur je eine Base des Tripletts ersetzt wird.

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The Effect of the Venom of the Digger Wasp Philanthus on the Fast and Slow Excitatory and the Inhibitory System in the Locust Muscle¹

Although many authors 2-11 have described the physiology of the fast and slow excitatory and the inhibitory neuromuscular system in insects, still very little is known about the pharmacology of these systems. A neuromuscular block of the fast system of the locust muscle can be caused by tryptamine and 5-hydroxytryptamine 12, gallamine 13, and the venom of Philanthus triangulum F. 13-15. Usherwood and Grundfest 9 showed that γ aminobutyric acid activates the inhibitory postsynaptic membrane in Romalea and Schistocerca. Although picrotoxin abolishes the inhibitory postsynaptic potentials and also the effect of γ -aminobutyric acid, it does not affect the excitatory postsynaptic potentials of the fast system 9.

In the present paper some effects of Philanthus venom on the fast and slow excitatory and on the inhibitory system of the locust jumping muscle will be described.

The wasps (Philanthus triangulum F.) originating from the South of France are reared in our laboratory by Dr. R. T. Simon Thomas 16. The method of extracting the venom as well as the preparation of the locust (Locusta migratoria migratorioides R. & F.) has been described in a preceding paper 13.

Intracellular action potentials were recorded from the upper part of the extensor muscle of the tibia of the metathoracic leg, using glass capillary microelectrodes filled with 3M KCl. The electrodes with a resistance between 20 and $40 M\Omega$ were connected to the input of a unity gain negative capacity preamplifier.

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