

Stimulating illusory own-body perceptions

The part of the brain that can induce out-of-body experiences has been located.

'Out-of-body' experiences (OBEs) are curious, usually brief sensations in which a person's consciousness seems to become detached from the body and take up a remote viewing position¹⁻³. Here we describe the repeated induction of this experience by focal electrical stimulation of the brain's right angular gyrus in a patient who was undergoing evaluation for epilepsy treatment. Stimulation at this site also elicited illusory transformations of the patient's arm and legs (complex somatosensory responses) and whole-body displacements (vestibular responses), indicating that out-of-body experiences may reflect a failure by the brain to integrate complex somatosensory and vestibular information¹⁻³.

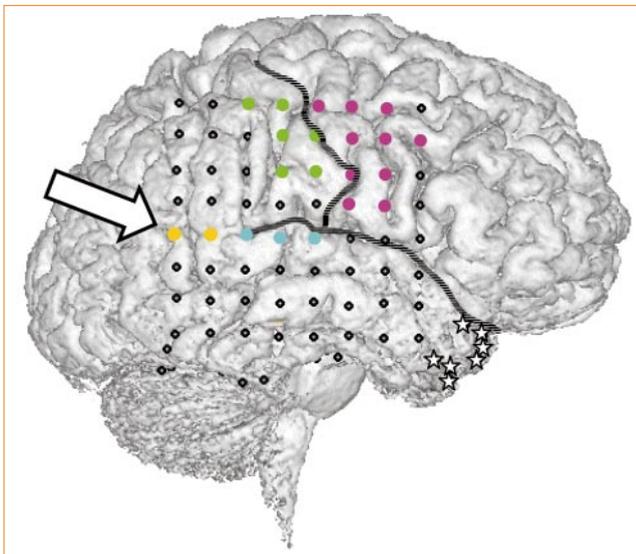
Our patient was a 43-year-old, right-handed woman who had suffered from complex partial seizures for 11 years; right temporal-lobe epilepsy was implicated. As magnetic-resonance imaging did not reveal any lesion, invasive monitoring was undertaken to localize the seizure focus precisely. Subdural electrodes were implanted to record seizures, and focal electrical stimulation was used to identify the vital cortex⁴.

Figure 1 shows the results of stimulation mapping and the electrode site on the right angular gyrus where stimulation repeatedly induced OBEs, as well as vestibular and complex somatosensory responses. Mapping of motor, somatosensory and auditory functions revealed no deviant brain pathology in this patient with respect to anatomical representations of cortical functions. The epileptic focus was located more than 5 cm anterior to the stimulation site, in the medial temporal lobe; electrical stimulation of this site did not induce OBEs, and these experiences were not part of the patient's habitual seizures.

Initial stimulations ($n = 3$; 2.0–3.0 mA) induced vestibular responses, in which the patient reported that she was "sinking into the bed" or "falling from a height". Increasing the current amplitude (3.5 mA) led to an OBE ("I see myself lying in bed, from above, but I only see my legs and lower trunk"). Two further stimulations induced the same sensation, which included an instantaneous feeling of "lightness" and "floating" about two metres above the bed, close to the ceiling.

The patient was then asked to watch her (real) legs during the electrical stimulation ($n = 2$; 4.0, 4.5 mA). As before, she was lying down (upper body supported at an angle of 45°, legs outstretched). This time, she reported seeing her legs "becoming shorter".

Figure 1 Three-dimensional surface reconstruction of the right hemisphere of the brain from magnetic-resonance imaging. Subdural electrodes were implanted in the brain of an epileptic patient undergoing presurgical evaluation; the locations at which focal electrical stimulation (ES) evoked behavioural responses are shown: magenta, motor; green, somatosensory cortex; turquoise, auditory cortex. Yellow, site at which out-of-body experience (OBE), body-part illusions and vestibular responses were induced (arrow). Stars indicate the epileptic focus in the medial temporal lobe. Informed consent was obtained from the patient and ES procedures conformed to the Declaration of Helsinki. Constant current (0.5–5.0 mA, 2-s train duration) was applied at 50 Hz in a bipolar manner through adjacent contacts⁴. Since undergoing a right anterior temporal lobectomy in 2000, the patient has been free of complex partial seizures.



If the patient's legs were bent before the stimulation (90° knee angle; $n = 2$; 4.0, 5.0 mA), she reported that her legs appeared to be moving quickly towards her face, and took evasive action.

When asked to look at her outstretched arms during the electrical stimulation ($n = 2$; 4.5, 5.0 mA), the patient felt as though her left arm was shortened; the right arm was unaffected. If both arms were in the same position but bent by 90° at the elbow, she felt that her left lower arm and hand were moving towards her face ($n = 2$; 4.5, 5.0 mA). When her eyes were shut, she felt that her upper body was moving towards her legs, which were stable ($n = 2$; 4.0, 5.0 mA).

These observations indicate that OBEs and complex somatosensory illusions can be artificially induced by electrical stimulation of the cortex. The association of these phenomena and their anatomical selectivity suggest that they have a common origin in body-related processing¹⁻³, an idea that is supported by the restriction of these visual experiences to the patient's own body.

During her OBE, the patient only 'saw' that part of her body that she also felt was modified during her body-transformation experiences. This contrasts with the non-corporeal visual hallucinations that are commonly induced by electrical stimulation at the parieto-temporal junction⁵. As suggested by previous neurological investigations on OBEs¹⁻³ and other body-cognition disorders⁶⁻⁸, the angular gyrus could be a

crucial node in a larger neural circuit that mediates complex own-body perception.

Out-of-body and body-transformation experiences are transitory and may disappear when a person attempts to inspect the illusory body or body part¹⁻³. Our findings suggest that changes in visual attention and/or current amplitude in the angular gyrus^{4,9} could bring about these phenomenological modifications.

Although we do not fully understand the neurological mechanism that causes OBEs, our results imply that vestibular processing² may be important. Although translocational vestibular responses were evoked initially without an OBE and can be produced in isolation⁴, vestibular sensations of levitation and lightness¹⁻³ accompanied OBEs in our patient. Also, the core region of the human vestibular cortex is situated close to the angular gyrus¹⁰. It is possible that the experience of dissociation of self from the body is a result of failure to integrate complex somatosensory and vestibular information.

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Eukaryotic evolution

Early origin of canonical introns

Spliceosomal introns, one of the hallmarks of eukaryotic genomes, were thought to have originated late in evolution^{1,2} and were assumed not to exist in eukaryotes that diverged early — until the discovery of a single intron with an aberrant splice boundary in the primitive ‘protozoan’ *Giardia*³. Here we describe introns from a close relative of *Giardia*,

Carpediemonas membranifera, that have boundary sequences of the normal eukaryotic type, indicating that canonical introns are likely to have arisen very early in eukaryotic evolution.

Carpediemonas membranifera is a poorly studied, free-living microbial eukaryote that is considered to be a relative of *Giardia* on the basis of its morphology⁴. Using the polymerase chain reaction (PCR) with *Carpediemonas* genomic DNA as template, we determined the partial sequences of two distinct carbamate kinase genes from this organism. In both genes, an insertion of 33 or 31 nucleotides interrupts the similar protein-coding sequence shared with carbamate kinase genes from other organisms (Fig. 1a). These insertions are bounded by guanine and thymine (GT) nucleotides at the 5’ end and adenine and guanine (AG) nucleotides at the 3’ end, which is a characteristic of most of the spliceosomal introns that interrupt protein-coding genes in other eukaryotes.

We used PCR with reverse transcription to recover the messenger RNA sequence of one of the two *Carpediemonas* carbamate kinase genes. This sequence lacks the insertion, which is presumably removed (spliced) from the messenger RNA before translation. We conclude that the insertions in the *Carpediemonas* carbamate kinase genes are canonical ‘GT...AG’ spliceosomal introns, albeit comparatively small ones.

To determine the evolutionary affinities of *Carpediemonas*, we used PCR to amplify near-complete sequences for two genes that encode cytosolic heat-shock protein 70 (Hsp70). We also sequenced a cloned Hsp70 gene from *Spironucleus barkhanus*, a very close relative of *Giardia*. Maximum likelihood analysis of Hsp70 proteins reveals a specific evolutionary relation between *Carpediemonas*, *Giardia* and *Spironucleus* (Fig. 1b); three other molecular markers also support this relationship⁵.

The single intron found in a *Giardia* gene has a non-canonical CT dinucleotide at its 5’ splicing boundary³, which could be interpreted as a ‘frozen’ primitive eukaryotic condition: canonical ‘GT...AG’

spliceosomal introns might then be a later innovation in more modern cells. Our results indicate that this is not the case, however, as canonical introns seem to be an ancestral feature of the larger evolutionary grouping that includes *Giardia* and *Carpediemonas*. The aberrant *Giardia* intron probably represents a lineage-specific (or intron-specific) secondary alteration of the 5’ splice boundary.

The extremely early divergence attributed to *Giardia* is based on the absence or aberration of many typical eukaryotic features, such as mitochondria and introns, and on its arguably deep-branching position in many phylogenetic trees^{6–9}. The grouping of *Giardia* with *Carpediemonas* (which, as well as canonical introns, has organelles that are probably derived from mitochondria⁴) weakens this argument for early divergence.

Respective of the true evolutionary position of *Giardia*, the only potentially ‘early’ eukaryotic group in which introns have not been found are the parabasalids, such as *Trichomonas*^{10,11}. *Trichomonas* is already known to possess some of the cellular machinery for intron splicing¹², however, and there is evidence to indicate that it is evolutionarily affiliated with *Giardia* and its relatives^{7,13} (and is slightly misplaced in many phylogenies, including that shown in Fig. 1b). An affiliation with *Giardia* implies a similar closeness to *Carpediemonas*, and it is likely that parabasalids have, or had, canonical introns. There is now every reason to assume that canonical introns were present in the most recent common ancestor of living eukaryotes.

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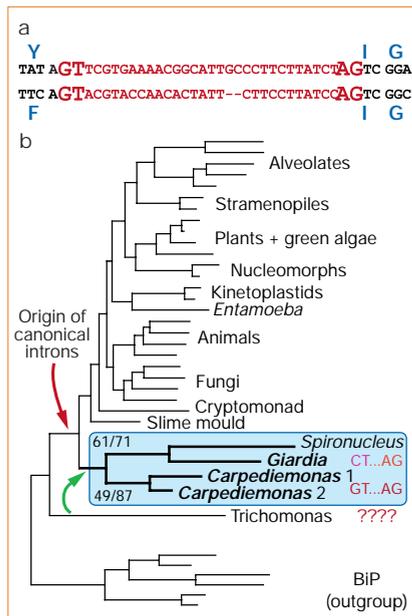


Figure 1 Introns and evolutionary affinities of *Carpediemonas*. **a**, Portions of two *Carpediemonas* carbamate kinase genes, showing intron sequences (red) interrupting the protein-coding sequence (in blue). The introns have canonical splice boundaries (GT...AG; large red type). **b**, Maximum-likelihood evolutionary tree of eukaryotic cytosolic Hsp70 proteins (‘T + invariable sites’ model). Endoplasmic-reticulum Hsp70 (‘BIP’) is used as an outgroup. The grouping of *Carpediemonas* with *Giardia* and *Spirionucleus* is shown in the blue box; statistical support (bootstrap percentages) for this grouping is assessed using likelihood (upper left of box), and likelihood distance (lower left). The higher percentages (right in each pair) apply when the outgroup is omitted. The basal placement of *Trichomonas* is weakly supported with likelihood (21%); green arrow shows a more plausible position on the basis of other evidence^{7,13}. The intron splice boundaries for the relevant groups and the origin of canonical introns are shown in red. New sequences have been deposited at GenBank under accession numbers AY131204–AY131209.

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