

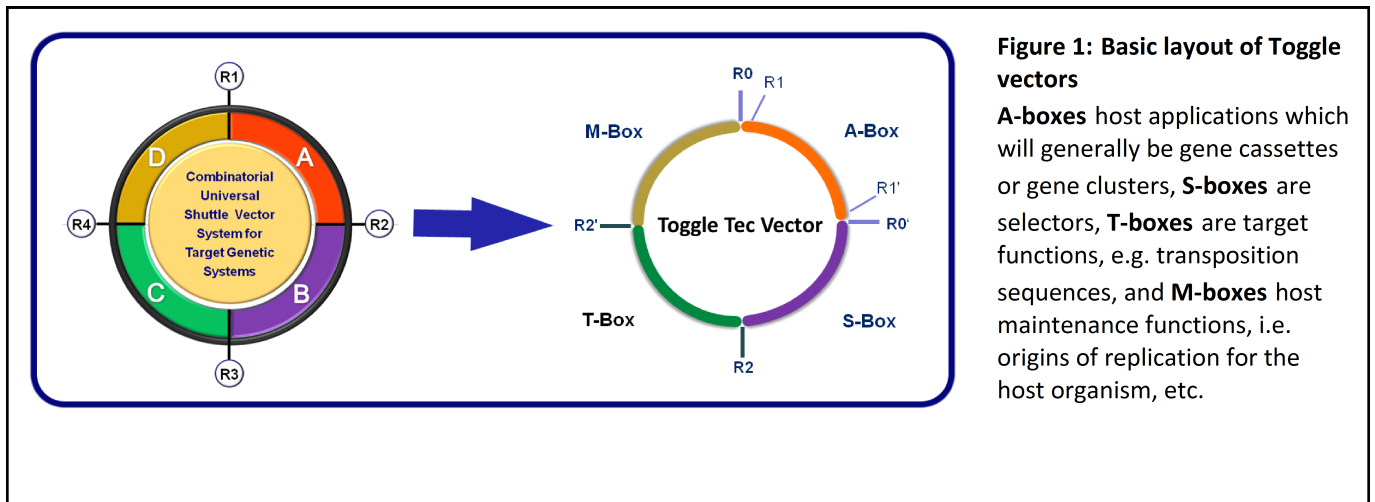
ACDC-SD and ToggleTec vectors: a new system for constructing modular multi-gene expression constructs

This newsletter describes how the assembly / disassembly cloning and substitution design (**ACDC-SD**) principle and a series of corresponding kits for different expression hosts will help you tackle such challenges in building and efficiently modifying such multi-gene expression assemblies effectively.

Organismal genes that encode **integrated protein machines** are frequently lumped into functional clusters, e.g. those involved in nonribosomal peptide synthesis (**NRPS**). The component proteins act in highly and tightly organized complexes to exert their physiological function.

The artificial integration of genetic function - such as production-optimized **educt-product** chains (e.g. in metabolic engineering) – at the functional level of biocatalysis is a challenge for synthetic biologists. The same holds true for probing and modifying structure-function relationships, e.g. when investigating intracellular or membrane-bound protein complexes that make up certain regulatory pathways.

ACDC-SD is fully compatible with and implemented in the **Toggle assembly system (ToggleTec vectors)**. Any gene to be inserted into the vector thus needs to adhere to a limited number of design principles to exploit the full potential of the system (fig. 2).



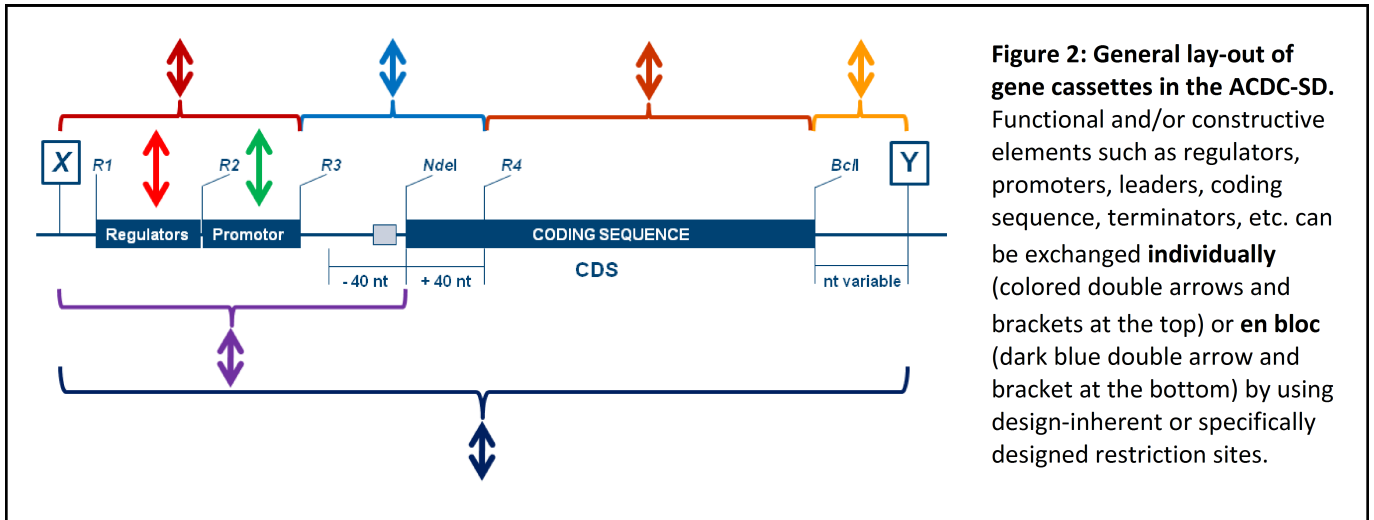
ATG offers ready-made systems for E.coli, insect and mammalian cells but can also assemble custom-made expression systems tailored to your host of interest.

Advantages of ATG's fully design-optimized ACDC-SD gene cassettes:

- Proven functional designs of gene clusters based on many years of expertise in **constructional synthetic biology**
- **Unified Design Concept** to realize genes, expression boxes/ cassettes, regulators etc.
- **Designs** incorporate a **de-assembly** option to regenerate individual expression boxes ? enables gene cluster revision and reorganization by **assembling** and/or **reusing** existing genetic elements

(fig. between sites X and Y)

- **Option for substitution cloning:** substitution of individual genetic elements or whole expression boxes in existing gene clusters (see fig. above)
- **Option for positional gene libraries** for the local evaluation of variants in a given gene-cluster construct



ACDC-S design allows you to use standardized expression boxes where all genetic elements can be recycled and exchanged between synthetic biologists who use the same standards for different organisms. Swapping ACDC-S elements between gene clusters and researchers who work on the same project is easy.

All systems support a diversity of assembly strategies for the development of higher-order integrated genetic functionality, e.g.

- **Ligation strategies** such as **DONOR/ ACCEPTOR**-based methods with alternating selector toggle assembly
- **Exonuclease cloning** ($5' \rightarrow 3'$, $3' \rightarrow 5'$)
- **Recombination-mediated assembly**, e.g. Cre/LoxP

Classical multiple cloning sites can be introduced into the systems as well.

ACDC-S is suited for these **applications:**

- **multi-gene artificial biochemical pathway systems** – in vivo e.g.
 - *anabolic pathways* – biologically active compounds, fine chemicals, building blocks, biofuels etc.
 - *catabolic pathways* – biorefineries, bioremediation etc.
- **multi-gene hetero-protein-complex systems** – in vivo
- multiple in vitro bio-catalyst co-expression systems (*flexiBAC*)
- other: specific functionally targeted multi-genetic constructive designs – in vivo
- multiple-resistance mechanisms in plants etc.

Order your **ACDC-S** design for preassembled gene clusters of your choice or the special kits that allow you to assemble your gene clusters according to the **ACDC-S** design.

ATG is *your* “**Synthetic Biology**” system provider that provides fully integrated molecular service packages to help you improve the quality of your research and to accelerate your developments.

Shape the future world of molecular design ... with ATG:biosynthetics!