Abstract. Classical cadherins form parallel cis-dimers that emanate from a single cell surface. It is thought that the cis-dimeric form is active in cell–cell adhesion, whereas cadherin monomers are likely to be inactive. Currently, cis-dimers have been shown to exist only between cadherins of the same type. Here, we show the specific formation of cis-heterodimers between N- and R-cadherins. E-cadherin cannot participate in these complexes. Cells coexpressing N- and R-cadherins show homophilic adhesion in which these proteins coassociate at cell–cell interfaces. We performed site-directed mutagenesis studies, the results of which support the strand dimer model for cis-dimerization. Furthermore, we show that when N- and R-cadherins are coexpressed in neurons in vitro, the two cadherins colocalize at certain neural synapses, implying biological relevance for these complexes. The present study provides a novel paradigm for cadherin interaction whereby selective cis-heterodimer formation may generate new functional units to mediate cell–cell adhesion.

Key words: cadherins • cell adhesion • heterophilic binding • synaptic adhesion • dimerization

Introduction

The cadherins are single-pass transmembrane proteins that mediate calcium-dependent cell–cell adhesion (Takeichi, 1991; Gumbiner, 1996). This superfamily consists of several subfamilies comprising the classical, desmosomal, and the CNR/protocadherins (Sano et al., 1993; Suzuki, 1996; Suzuki et al., 1997; Takeichi et al., 1997; Kohmura et al., 1998; Wu and M aniatis, 1999). Classical cadherins (for example, N-, E-, and R-) share a common primary structure with five tandemly repeated extracellular domains of ~110 amino acids each, a transmembrane segment, and a short cytoplasmic domain (Hatta et al., 1988; Matsunami et al., 1993; Shapiro and Colman, 1998). The cytoplasmic domain of the classical cadherins interacts with a complex that includes a- and b-catenins, and plakoglobin, all of which function in part to mediate connections between the cadherins and the cytoskeleton (Ozawa et al., 1989, 1990; Reynolds et al., 1992; Kemler, 1993; Shibamoto et al., 1995; Yap et al., 1998).

The first extracellular domain (EC1)1 governs the binding specificity of the cadherins. The best evidence for this comes from experiments which show that specificities of cell adhesion between P- and E-cadherins could be switched when their NH2-terminal regions were exchanged (Nose et al., 1990). Furthermore, site-directed mutations in the EC1 domain can significantly alter or abolish adhesion (Tamura et al., 1998).

Crystal structures of the EC1 domain from N-cadherin suggested new ideas about the adhesive interface between cadherins. Two separate interfaces, the strand dimer (which mediates parallel or cis-dimerization) and the adhesion dimer, are thought to contribute to N-cadherin adhesive function (Shapiro et al., 1995). The strand dimer is formed by lateral association between the side chain of tryptophan 2 (Trp-2) and a pocket in the hydrophobic core of the partner molecule. Mutagenesis studies have shown that Trp-2, and also its acceptor pocket, are crucial for adhesion in both N- and E-cadherins (Tamura et al., 1998). The necessity for lateral cis-dimers is supported by experiments with a recombinant C-cadherin extracellular segment (Brieher et al., 1996), showing that only the dimeric form mediates strong adhesion. Furthermore, artificially induced multimerization of a chimeric tailless C-cadherin could strengthen adhesion in cells expressing this molecule (Yap et al., 1997). Thus, it appears likely that cis-dimers are a fundamental unit for cadherin function, at least for the classical cadherins. More recent crystallographic evidence has been presented for both N- and...
E-cadherins, however, which either shows Trp-2 disordered (Nagar et al., 1996; Tamura et al., 1998) or inserted into a pocket in its own hydrophobic core (Pertz et al., 1999), rather than that of a dimer partner. This evidence does not necessarily dispute the strand dimer model. By analogy, enzymes can have active forms and inactive forms, and it is well known that much of the functional diversity of proteins accrues due to their ability to adopt alternate conformations. Therefore, it seems possible that the strand dimer and other conformations of Trp-2 and other nearby regions could reflect different natural states of cadherins.

In nervous tissue, classical cadherins are critical for formation of brain nuclei (Beesley et al., 1995; Redies and Takeichi, 1996; U chida et al., 1996; Colman, 1997; Redies, 1997; A rndt et al., 1998; U emura, 1998) and establishment of neuronal connectivity, including synaptic target recognition (Fannon and Colman, 1996; U chida et al., 1996). It is known that dozens of cadherins are expressed in brain (Suzuki et al., 1991, 1997; Sano et al., 1993; Suzuki, 1996), and at least some are coexpressed in the same neuron (K ohmura et al., 1998; W ohrn et al., 1999). A though N- and R-cadherins are related to each other, the expression patterns of N- and R-cadherins in the central nervous system differ substantially from one another (Redies et al., 1993; Ganzler and Redies, 1995; M atsunami and Takeichi, 1995; Redies, 1997; W ohrn et al., 1998). Partial overlap in expression of N- and R-cadherins in the same cell groups has been demonstrated (I nuzuka et al., 1991; Redies et al., 1993; Redies, 1997; W ohrn et al., 1999). Therefore, various interactions between N- and R-cadherins, and perhaps other cadherins, may exist to maintain the integrity of nervous and other tissues.

To investigate the possibility of heterodimerization between different cadherins, we expressed N-, E-, and R-cadherins in L cells and established stable single and double cadherin expressors. We chose to study N- and R-cadherins because regions of N-cadherin expression in the nervous system often abut regions of R-cadherin expression (I nuzuka et al., 1991; R edies et al., 1993; R edies, 1995; W ohrn et al., 1998, 1999). In contrast, the expression pattern of E-cadherin shows no such correlation (M atsunami and Takeichi, 1995; Fannon and Colman, 1996). We show here that N- and R-cadherins can form stable cis-heterodimers, whereas E-cadherin cannot dimerize with either of these molecules. Site-directed mutagenesis experiments suggest that N/R cis-dimers are likely to adopt the strand dimer configuration. These heterodimers can interact with appropriate cadherins from opposing cells to mediate functional cell–cell adhesion. The correlated expression, both in time and space, observed for N- and R-cadherins suggests that the functional cis-heterodimers between N- and R-cadherins reported here may reflect their natural interaction in vivo. It is of interest that the formation of parallel dimers allows for cis-heterodimer formation, while maintaining homophilic adhesive interactions across the opposed bilayers.

Materials and Methods

Antibodies

Rat mAbs against the extracellular domain of mouse N-cadherin (M NCD2) and the extracellular domain of mouse R-cadherin (M RCD5) were kindly provided by Prof. M. Takeichi (K yoto U niversity, j apan; M atsunami and Takeichi, 1995). Polyclonal antibodies against the EC1 domain of N-cadherin (anti-NEC1) and the ECS domain of E-cadherin (anti-ECS5) were generated in rabbit and guinea pig, respectively (F annon and Colman, 1996). Rabbit polyclonal antibody against GFP was purchased from Clontech Laboratories. M ouse mAbs against myc tag and rabbit polyclonal antibodies against α- and β-catenin were purchased from Sigma Chemical Co.

cDNA Constructs
cDNAs encoding full-length N- and E-cadherin were isolated by screening a mouse brain cDNA library. The coding regions were sequenced and were identical to the published N- and E-cadherin sequences (Nagafuchi et al., 1987; M iyatani et al., 1989). R-cadherin cDNA in Bluescript was kindly provided by Prof. M. Takeichi (M atsunami et al., 1993). The N-, E-, and R-cadherin coding regions were each inserted into the pCXN2 expression vector for protein expression in L cells.

For immunoprecipitation, R-cadherin was fused with GFP (green fluorescence protein) at its COOH terminus and N-cadherin with 6 copies of the myc epitope. A full-length R-cadherin–GFP and N-cadherin–myc was inserted into the pCXN2 vector. The site-directed mutation in EC1 domain of N-cadherin, Trp2→Ala (W2A), which causes dissociation of N-cadherin dimer and loss of adhesive activity, was generated as described before (Tamura et al., 1998). A deletion mutation of the N-cadherin EC1 domain was generated by removing the coding region from 710-1,256 by EcoNI and ligated. These mutations did not change the reading frame. To generate a chimeric construct ENEC1, the E-cadherin EC1 domain was substituted with the N-cadherin EC1 domain. For this procedure, a 1.5-kb fragment of E-cadherin containing signal peptide and mature peptide EC1–EC3 region was cut out and subcloned into XbaI and XhoI site of Bluescript vector. A PCR fragment containing a new Tth111I restriction site in E-cadherin was cloned into XhoI and ligated. The whole chimeric cadherin cDNA was cloned into the pCXN2 vector. A PCR products and ligation sites were sequenced using the ABI computerized automated DNA sequencing.

Cell Culture and Transfection

L cells were cultured in a humidified atmosphere of 10% CO2 and in 90% DMEM containing 10% FBS. Neurons were prepared from hippocampi of embryonic day 18 Sprague Dawley rats as described previously (Benson and Colman, 1998). In brief, neurons were dissociated by surgical trituration through a Pasteur pipette. Neurons were plated at a density of 3,600 cell/cm2 on poly-l-lysine-coated coverslips in MEM containing 10% horse serum. After 4 h, when neurons had attached, coverslips were transferred to dishes containing a monolayer of cortical astroglia, where they were maintained for up to 2 wk in MEM containing N2 supplement, sodium pyruvate (1 mM), and ovalbumin (0.1%). To obtain single or double transfectants, the cells were transfected by using Superfect (Q iagen Inc.), and then cultured in selective medium containing 800 μg/ml of G418 ( Gibco BRL). Colonies were isolated and examined for N-, R-, and E-cadherin expression by immunofluorescence and positive cells were expanded and used for further studies.

Immunocytochemistry

Cells were fixed in 4% paraformaldehyde, delipidated in 100% methanol, permeabilized with 0.1% Triton X-100, and blocked with 5% normal goat serum in PBS. A 20-min incubation with primary antibodies, cells were then incubated with fluorescent-conjugated secondary antibodies (Jackson Immunoresearch Laboratories) at room temperature for 30 min. Coverslips were then mounted and examined by confocal laser microscopy.

Aggregation Assays

Monolayer cultures were treated with 0.01% trypsin in HCMF (Hepes-buffered Ca2+- and Mg2+-free Hanks’ Solution) supplemented with 1 mM MgCl2. Neurons were treated with 0.01% trypsin in HCMF (Hepes-buffered Ca2+ and Mg2+-free Hanks’ Solution) supplemented with 1 mM MgCl2.
CaCl$_2$ for 30 min at 37°C. The trypsinized cells were washed gently in HCMF containing calcium and 1% BSA at 4°C. This procedure can dissociate cell layers into single cells, leaving cadherins intact on the cell surface. After the cells were thoroughly dissociated, 5 x 10$^5$ cells per well were transferred to 24-well dishes for a final volume of 0.5 ml HCMF containing 1% BSA with or without 1 mM Ca$^{2+}$. The plates were rotated at 80 rpm at 37°C for 45 min.

For mixed aggregation analysis, cells expressing different cadherin constructs were labeled with different lipophilic dyes before mixing. We used DiI (1,1-dioctadecyl-3,3,3,3-tetramethylindocarbocyanine) and DiO (3,3-diododecylxocarbocyanine perchlorate), each purchased from Molecular Probes. Stock solutions of DiI were made by dissolving 2.5 mg DiI in 1 ml of 100% ethanol, and stocks of DiO were made by dissolving 2.5 mg DiO in 1 ml of 90% ethanol, 10% dimethylsulfoxide. These stock solutions were sonicated and filtered before use. To label cells with these dyes, they were incubated for 8 h at 37°C in serum-containing DME at final concentrations of 15 mg/ml and 10 mg/ml for DiI and DiO, respectively. The cells were washed extensively with HCMF containing calcium to prevent cross contamination of the dyes. After single cell suspensions were obtained as described above, 2.5 x 10$^5$ cells per well of each of two types were transferred to a 24-well dish. After rotating at 80 rpm at 37°C for 45 min, 50 μl of the fixed aggregates were removed, placed on a slide, and covered with a coverslip and examined by confocal microscopy.

**Immunoblotting and Immunoprecipitation**

For immunoblotting and immunoprecipitation experiments, 5 x 10$^6$ cells were cultured in 100-mm tissue culture dishes at 37°C for 3 d. In coculture experiments, 5 x 10$^6$ cells expressing N-cadherin wild-type and R-cadherin-GFP mixed in a ratio 1:1 were cultured in a 100-mm dish for 24 h. The confluent monolayer was then washed with PBS, scraped off the dish, pelleted for 5 min at 1,000 rpm and extracted in 1.0 ml of lysis buffer (150 mM NaCl, 20 mM Tris-HCl, pH 7.5, 1% NP-40, 1% Triton X-100, Ca$^{2+}$ free leupeptin 0.5 μg/ml, pefabloc 0.1 mM, aprotinin 1 μg/ml) for 30 min. After cell lysis, the samples were centrifuged for 20 min at 15,000 rpm and the supernatants were separated. Protein concentration was determined with the BCA protein assay (Pierce Chemical Co.) and samples were run on 6% SDS-PAGE, transferred to nitrocellulose, membrane-blocked with 5% milk protein, and incubated overnight with primary antibodies. After secondary antibody incubation and routine washing, blots were developed.
with the chemiluminescence system (New England Nuclear Life Science Products). For immunoprecipitation, the supernatants were incubated with specific antibody (3 μl/sample) for 1 h at 4°C. The antigen–IgG complexes were precipitated by sequential incubation with protein-A-Sepharose 4B. The immunoprecipitates were washed extensively, and boiled with SDS-sample buffer, and the proteins were detected by immunoblotting with specific antibodies.

For N-cadherin or GFP immunodepletion experiments, 1 ml cell lysates were incubated with 20 μl of N-cadherin or 10 μl GFP antibodies. The antibodies with complexes were then removed by anti-rabbit IgG-agarose (Sigma Chemical Co.). 30 μl of each protein sample from the homogenates was subjected to SDS-PAGE, and the separated proteins were electrophoretically transferred to a nitrocellulose membrane sheet. The membrane was processed to detect protein with specific antibodies.

**Figure 2.** R-/N-cadherin, when coexpressed in L cells, forms heterocomplexes. A, Immunostaining of double transfectants showed that different cadherins were coexpressed in L cells. Bar, 10 μm. B, Lysates from L_R, L_N, and L_R/N cells were immunoprecipitated with GFP antibody (IP: GFP) and detected by N- and R-cadherin and α- and β-catenin antibodies. N-cadherin was found to be coprecipitated with R-cadherin in L_R/N cells. C, Lysates or immunoprecipitates from L_E, L_R/E_1, and L_R/E_2 cell lines were detected by E- and R-cadherin and α- and β-catenin antibodies.
Results

**R- and N-Cadherins Form Heterophilic Cell–cell Interfaces**

To evaluate and compare the adhesive properties of N-, R-, and E-cadherins, we generated stable L cell lines expressing these cadherins (L<sub>R</sub>, L<sub>N</sub>, and L<sub>E</sub>). Aggregation assays were performed to verify the adhesive properties of these cell lines. Stable transfectants of each cadherin showed rapid calcium-dependent formation of large aggregates.

To test the binding specificity, we performed mixed aggregation assays with DiI (red) or DiO (green)-labeled cell lines. When L<sub>R</sub> cells were mixed with L<sub>N</sub> cells, heterotypic aggregates were produced, although the individual cell types still remained segregated (Fig. 1 a). In contrast, mixed assays of L<sub>E</sub> cells with either L<sub>N</sub> or L<sub>R</sub> cells showed that L<sub>E</sub> cell aggregates remained entirely segregated from L<sub>R</sub> or L<sub>N</sub> cell aggregates (Fig. 1, b and c). These results were consistent with previous reports (Matsunami et al., 1993) and suggest that the heterophilic interaction of R- and N-cadherins may be weaker than their respective homophilic interactions. Similarly, this interaction also can be shown in L<sub>R</sub> and L<sub>N</sub> cell cocultures by immunostaining using N-cadherin antibodies and R-cadherin–GFP (Fig. 1, d–f). Both types of cadherins were present at the junctions between the two subpopulations, even when the two cell types were mixed randomly before plating on the dish. Complexes of R- and N-cadherins, however, could not be detected by coimmunoprecipitation using GFP antibody (see Fig. 3 A). These results indicate that N- and R-cadherins can form heteroadhesive clusters between adjacent cells, but these trans-complexes are too weak to maintain under condition used for routine immunoprecipitation.

Cocultures of E- and R-cadherin expressors, and E- and N-cadherin expressors, were also examined (Fig. 1, g–l). Immunolocalization of the respective cadherins showed that N- and R-cadherins each formed cell junctions that were clearly separate from E-cadherin-mediated junctions. This is in contrast to N/R cocultures which clearly show regions of heterotypic adhesion between N- and R-cadherins. This leads us to suggest that the differential adhesive properties of these molecules are important factors in controlling their localization at cell junctions.

**R- and N-Cadherins Form Stable Heterodimers in Coexpressing Cell Lines**

We investigated whether different cadherins could form cis-heterodimers by using cell lines coexpressing R- and N-cadherins or R- and E-cadherins (L<sub>R/N</sub>, L<sub>R/E</sub>; Fig. 2 A). We used an R-cadherin fused with GFP to enable detection in coimmunoprecipitation experiments. Total cell lysates from L<sub>N</sub>, L<sub>R</sub>, and L<sub>R/N</sub> were immunoprecipitated with GFP antibody and detected by Western blotting with N-, R-, or E-cadherin antibodies. The R-cadherin–GFP fusion protein was able to coprecipitate with N-cadherin from the L<sub>R/N</sub> cell lysate (Fig. 2 B). In contrast, E-cadherin was not detected in immunoprecipitates from L<sub>R/E</sub> cell lines (Fig. 2 C). These two proteins, however, accumulate at cell membranes (Fig. 2 A). These data indicate the absence of interaction between R- and E-cadherins. In both L<sub>R/N</sub> and...
L<sub>R</sub>/E cell lines, cadherins were associated with α- and β-catenin and these proteins could be precipitated together with GFP antibody, suggesting that catenins do not control the heterointeractions between these cadherins (Fig. 2, B and C).

Heterophilic interaction in a calcium-dependent manner has been shown between L<sub>R</sub> and L<sub>N</sub> cells, both in our experiments and in those previously reported (Matsunami et al., 1993). Therefore, we used a calcium chelator (EGTA) to disrupt the adhesive interaction, so that we could determine whether the R-/N-cadherin complexes obtained by immunoprecipitation originated from the same or opposing cells. We made lysates from L<sub>N</sub> and L<sub>R</sub> cocultures treated with or without EGTA and performed immunoprecipitations with GFP antibody. R-cadherin–GFP did not coprecipitate N-cadherin in mixed monoexpressing L<sub>N</sub> and L<sub>R</sub> cocultures, either in the presence or absence of EGTA (Fig. 2, B and C). These results imply that the trans-heterophilic adhesive complex between R- and N-cadherin may be considerably weaker than the respective homophilic counterparts and undetectable in our assay system. In contrast, when immunoprecipitations were performed on lysates from the double-transfectant, L<sub>R/N</sub>, in the presence of EGTA (~20 mM), R-cadherin–GFP coprecipitated N-cadherin equally as well as in the presence of Cal<sup>2+</sup> (Fig. 3 B). Taken together, our data indicate that lateral dimerization accounts exclusively for the interaction between R- and N-cadherins observed in the coprecipitation experiments, and that the maintenance of this interaction is independent of Cal<sup>2+</sup>. Similar data have shown that lateral dimerization of E-cadherin is also independent of Cal<sup>2+</sup> (Chitaev and Troyanovsky, 1998).

We took great care to ensure that lateral dimerization between R- and N-cadherins could form only in cotransfected cells, and was not an artifact created in the lysate when the two proteins were initially mixed together. The lysates from L<sub>N</sub> and L<sub>R</sub> were pooled, incubated with or without Cal<sup>2+</sup> for 1 h, and subjected to immunoprecipitation with GFP antibody. R-/N-cadherin complexes were detected only in cells in which these proteins were coexpressed and not when individual L<sub>R</sub> and L<sub>N</sub> lysates were mixed (Fig. 3 C). Furthermore, when the L<sub>R/N</sub> cell lysate was depleted with GFP and NEC1 antibodies, although the protein amount of R- and N-cadherin was reduced, both cadherins were still detectable (Fig. 3 D). Thus, cells stably transfected with R- and N-cadherin probably contain three association forms: R-/N-, N-/N-, and R-/R-cadherin dimers. The existence of monomeric forms also cannot be ruled out.

**Tryptophan 2 of N-Cadherin Is Essential for Heterodimerization of R- and N-Cadherins**

Previous studies demonstrated that Trp-2 in the N-cadherin EC1 domain is critical for strand dimer formation (Shapiro et al., 1995; Shapiro and Colman, 1998; Tamura et al., 1998). To evaluate whether this model is relevant to R-/N-cadherin heterodimer formation, several constructs were produced. These included: N-cadherin lacking the EC1 domain (N<sub>ΔEC1</sub>); Trp-2 mutated to Ala in N-cadherin (NW2A); and a chimeric molecule of E-cadherin whose EC1 domain was replaced by the N-cadherin EC1 domain (ENEC1; Fig. 4 A). R-cadherin–GFP and mutated N-cadherin were cotransfected into L cells to generate coexpressing cell lines, L<sub>R/N</sub>, L<sub>R/N,NΔEC1</sub>, and L<sub>R/N,NW2A</sub>, and immunoprecipitations were performed using GFP antibody. Either deletion of the EC1 domain or the NW2A mutation completely abolished the immunoprecipitation of N-cadherin with R-cadherin (Fig. 4, B and C). Therefore, the EC1 domain, and particularly Trp-2, is required for R-/N-cadherin heterodimer formation.

**The Specificity of Heterodimerization Is Localized to the NH<sub>2</sub>-terminal Domain**

Switching the cell-to-cell adhesive specificity of cadherins by EC1 domain exchange was demonstrated previously (Nose et al., 1990). To test whether the specificity of cis-dimer formation might also be encoded in the EC1 domain, we engineered a chimeric protein, ENEC1, formed by replacing the EC1 domain of E-cadherin by the EC1 domain (ENEC1; Fig. 4 A). R-cadherin–GFP and mutated N-cadherin were cotransfected into L cells to generate coexpressing cell lines, L<sub>R/N,NΔEC1</sub> and L<sub>R/N,NW2A</sub>, and immunoprecipitations were performed using GFP antibody. Either deletion of the EC1 domain or the NW2A mutation completely abolished the immunoprecipitation of N-cadherin with R-cadherin (Fig. 4, B and C). Therefore, the EC1 domain, and particularly Trp-2, is required for R-/N-cadherin heterodimer formation.
Figure 5. Binding specificity of ENEC1 and heterointeraction between R-cadherin and ENEC1. (A) L\textsubscript{ENECE1} or L\textsubscript{R/ENECE1} was immunostained by NEC1 and EEC5 antibodies, showing that chimeric E-cadherin and R-cadherin colocalized at cell–cell contacts. (B) L\textsubscript{ENECE1} cells coaggregated with L\textsubscript{N} cells, but aggregated to a lesser extent with L\textsubscript{E} and L\textsubscript{R} cells. L\textsubscript{ENECE1} was labeled with DiO (green), L\textsubscript{N}, L\textsubscript{E} and L\textsubscript{R} cells with DiI (red). (C) Immunoprecipitation of L\textsubscript{R/N}, L\textsubscript{R/ENECE1}, and L\textsubscript{R/E} cell lines with GFP antibody. Precipitates were detected with anti-EEC5, N-cadherin, R-cadherin, and α- and β-catenin, showing that ENEC1-cadherin coprecipitated with R-cadherin. Bars, 25 μm.
domain of N-cadherin. We tested the adhesive and lateral interaction properties of this chimeric molecule in L cells (L_{ENEC1}). Immunocytochemical analyses showed that the chimeric cadherin protein was expressed on the cell surface and was concentrated at cell-cell boundaries (Fig. 5 A). L_{ENEC1} cell lines exhibited strong self-aggregation. These cells also formed aggregates with L_N cells, but showed far weaker aggregation with L_E or L_R cells (Fig. 5 B), indicating that ENEC1 was a functional artificial cadherin with adhesive specificity appropriate for the N-cadherin type. To determine whether ENEC1 interacted with R-cadherin, we established L_{R,ENEC1} coexpressors by cotransfecting R-cadherin-GFP and ENEC1 into L cells. The R-cadherin and ENEC1-chimeric proteins accumulate at sites of cell-cell contacts by immunostaining with E-cadherin antibody and R-cadherin-GFP fluorescence (Fig. 5 A). Using L_{R,N} and L_{R,E} cell lines as positive and negative controls, L_{R,ENEC1} cell lysates were treated with GFP antibody and the immunoprecipitates were probed with EEC5, NEC1, and RCD5 antibodies. These experi-
ments (Fig. 5 C) showed that ENEC1 could be precipitated together with R-cadherin-GFP, suggesting that heterodimers were formed between R-cadherin and ENEC1. This suggests that the EC1 domain of classic cadherins plays an important role in determining the specificity of cis-heterodimer formation.

The R/N Heterodimer Constitutes a Functional Unit for Cell–Cell Adhesion

Can coexpressing L_{R,N} cell lines mediate functional cell adhesion? And, if so, are R/N heterodimers involved in this process? To answer these questions, we first performed aggregation assays between different cadherin expressing cell lines. L_{R,N} cells aggregated strongly with their own type and exhibited high affinity with both L_{N} and L_{R} cell lines, but segregated from L_{E} cells (Fig. 6 A), suggesting that L_{R,N} cell lines hold broad-range adhesive specificity. Second, we generated another cell line, L_{N/m}, which is a chimera of N-cadherin fused with a myc epitope tag, and cocultured these cells with L_{R,N} and L_{R/N/W2A} cells. The cell lysates were immunoprecipitated with either myc or GFP antibody. These experiments (Fig. 6 B) showed that R-/N-cadherin heterodimers could be coprecipitated with N-cadherin–myc in the L_{R,N} and L_{N/m} cocultures, but not in the L_{R,N/W2A} and L_{N/m} coculture system, implying that the R-/N-cadherin heterodimer from one cell was functionally associated with N-cadherin from the opposing cell. This is notably different from the observation that R-cadherin–GFP cannot precipitate N-cadherin from an opposing cell, suggesting that the N/N association is stronger than the N/R association (see Fig. 8 C). We have performed experiments which showed that the W2A mutant of N-cadherin is a simple loss of function mutant, with no discernable dominant-negative effects (Fig. 6 A, e–h). This indicates that cells cotransfected with an adhesive cadherin and the NW2A mutant cadherin exhibit adhesive behavior identical to that of the adhesive active cadherin.

To investigate whether the association between N-cadherin–myc and R-/N-cadherin heterodimers depend on cell–cell contacts, the cocultured cells were incubated for 10 min with different concentrations of EGTA before lysis. Here, data showed that N-cadherin–myc interacted with R-/N-cadherin heterodimers in a calcium-dependent manner. This suggests that the calcium-dependent association between N-cadherin–myc and R-/N-cadherin heterodimers is necessary for functional cell adhesion. The distribution of R- and N-cadherins after expression in hippocampal neurons. Cell surface labeling of R-cadherin–GFP (A and a), N-cadherin–myc (B and b), and synaptophysin (C and c) was present at synaptic sites. Localization of R- and N-cadherins and synaptophysin is shown in D and d. Bar, 5 μm.
manner. Dimers between cells dissociated from each other in high concentrations of EGTA (Fig. 6, C and D), whereas R-/N-cadherin cis-heterodimers were stable to EGTA treatment (Fig. 6 D). These data reveal that sensitivity to EGTA treatment is different between cis-heterodimers and the cell-to-cell adhesion dimers.

Overexpressed R- and N-Cadherins Colocalize at Synaptic Junctions

Overlapping expression of R- and N-cadherin in the same neuron or fiber fascicle has been observed in the CNS (Redies, 1997; Wohrn et al., 1999). Whether both cadherins can be present at the same synaptic junction is still unclear. We coexpressed R-cadherin-GFP and N-cadherin-myc in cultured hippocampal neurons. After a 72-h cotransfection, the neurons were fixed and stained using epitope tag antibodies. The results showed that R- and N-cadherins colocalize at the same synaptic junctions (Fig. 7, A–D, a–d). This colocalization suggests that the associations we observed between R- and N-cadherins in L cells may reflect similar interactions in neurons, where they may naturally be found together.

Discussion

The most important findings presented here are: 1, in cotransfected cell lines, R- and N-cadherins form stable cis-heterodimers; 2, Trp-2 in the EC1 domain of N-cadherin is required for heterodimer formation, suggesting association in a strand dimer conformation; 3, the heterodimer is a functional unit for mediating cell-cell adhesion; and 4, the cis-heterodimer and the adhesive dimer exhibit different requirements for Ca²⁺, implying that different interfaces mediate these interactions. These observations suggest that selective formation of strand heterodimers between different cadherins may play an important role in specific cell adhesion, and present the possibility that cis-heterodimers may be functionally relevant in vivo. While the data presented here does not prove that the oligomers identified by immunoprecipitation correspond to dimers, the circumstantial case is quite strong. The ability to abrogate these oligomers by specific mutation of Trp-2, the central structural residue in the crystallographically observed strand dimer, is strong supportive evidence.

In addition to the data presented here and the crystallographic data, dimeric interaction near the NH₂-terminal domain of E-cadherin was also observed by EM (Tomschy et al., 1996). These studies showed that the adhesive interaction between NH₂-terminal domains occurs by a two-step mechanism: antiparallel pairs from different molecules form a complex only after parallel pair formation. X-ray crystallographic studies of the NH₂-terminal of the E-cadherin 1 and 2 domains suggested a different parallel dimer (Nagar et al., 1996). Experiments with a recombinant C-cadherin extracellular segment (CEC1-5) also revealed the existence of cis-dimeric interactions. Dimers were observed by chemical cross-linking, both in the presence and absence of calcium, and it was also shown that the CEC1-5 dimers had greater homophilic binding activity than monomers (Brieher et al., 1996). Consistent with this, a tailless E-cadherin can be cross-linked as a functional dimer on the cell surface (Ozawa and Kemler, 1998). Chitaev and Troyanovsky (1998) showed that E-cadherin formed stable lateral dimers in epithelial junctions. Our present results provide the first evidence for parallel heterodimer formation between R- and N-cadherin, and suggest that heterodimer formation is specific to cadherin type. Furthermore, since R-cadherin makes heterodimers with a recombinant E-cadherin molecule whose EC1 domain was replaced by the N-cadherin EC1 domain, we conclude that the EC1 domain contributes to the recognition of parallel partner molecules, and that it controls the specificity of heterodimer formation.

In the current work, we show that mutating Trp-2 in N-cadherin can destroy heterodimer formation between R- and N-cadherin. For E-cadherin, the double mutation (Trp-2→A/aVal3→Gly) within the EC1 domain completely abolished dimerization, suggesting that dimer formation in E-cadherin was also driven by Trp-2 and hydrophobic core interaction (Chitaev and Troyanovsky, 1998), as is the case for N-cadherin homodimers. This region may therefore be critical for both homo- and heterodimerization. It is difficult to predict the structural determinants of cis-dimer specificity based on the one existing crystal
structure. While it appears that Trp-2, which is entirely conserved throughout the classical cadherins, is the central residue in the interface, other regions may function in determining specificity. We have not attempted to model these interactions due to the highly speculative nature of such studies. We do note, however, that the E C1 domain of N- and R-cadherins are far more similar (73% identity) than that of N- and E-cadherins (61%) or R- and E-cadherins (54%).

Our investigation raises the question of whether cadherin cis-heterodimers play a natural functional role in cell adhesion. We have shown that L\textsubscript{IN} cells have broad-range adhesive specificity and high affinity with both L\textsubscript{R} or L\textsubscript{N} cells. The R-/N-cadherin heterodimer forms adhesive complexes with N-cadherin from opposing cells, revealing that the heterodimer is a functional cell–cell adhesive unit. Furthermore, N- and R-cadherins colocalize on neuronal synapses on the same cell surface, suggesting the likelihood that they naturally form heterodimers. A though the exact nature of cadherin adhesive contact is still unknown, strand dimer formation seems to be a critical step for subsequent adhesive dimer formation, and potentially for the formation of zipper-like adhesive structures proposed previously (Shapiro et al., 1995). Homophilic and heterophilic binding (Fig. 8 A–C) were thought to mediate strong and weak adhesion for different processes in central nervous system development, such as neurite outgrowth and synaptic junction formation (Colman, 1997; Tanaka et al., 2000). Cis-heterodimeric interactions (Fig. 8 C) between different cadherins may be a biologically relevant mechanism in cell–cell adhesion, which could play an important role in controlling different states of adhesiveness between cells.

It is generally believed that the generation of adhesive specificities depends on amino acid differences in E C1 of a classical cadherin. The conservation of the Trp-2, as well as those residues forming the hydrophobic pocket that it inserts into, suggests the possibility that heterodimer formation might be feasible in those cells where more than one cadherin is expressed. Heterodimer formation between cadherins expressed by the same cell could open up interesting evolutionary possibilities for the testing of new adhesive specificities through gene duplication and natural selection, while at the same time preserving one member of the dimer pair in an unaltered functional homophilic adhesive bond. Furthermore, the possibility of titration of adhesive strength using different combinations of strong and weak cadherins would provide a way for cells to exploit a wide spectrum of adhesive responses, and even radically change their intercellular associations and affinities.

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